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score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Direct Submission
Submitted (14-MAY-1996) Dunn-Walters D.K., UMDS
Dept of Histopathology, Lambeth Palace Road, Lo
2 (bases 1 to 286)
Dunn-Walters, D.K. and Spencer, J.
Origins of lamina propria plasma cells
Proc. Ninth Int. Congress for Mucosal Immunol.
Location/Qualifiers
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M.vison mRNA for ryanodine r
Y07749
Y07749.1 GI:1526614
ryanodine receptor 3; RYR3 g
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DVLGAEKSQIAFPKMVASCCRFLCYFCRISRQNQKAMFEHLSYLLENSSVGLASPSMR
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NRLCFLEPTSEAKYVPPDLCVCNFVLEQSLSVRALQEMLANTGEACWWTIHPASKQRS
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FEFEVVTGGDMRVGWARPGCRPDIELGADDQAFVFEGSRGQRWHQGSGYFGRTWQPGD
VVGCMINLDDASMIFTLNGELLITNKGSELAFADYEIENGFVPICSLGLSQIGRMNLG
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FKTLTEKEKETYRWPARESLKTMLAVGWSVERTKEGEALVQQRENEKLRSVSQASQGN
SYSPAPLDLSNVVLSRELQGMVEVVAENYHNIWAKKKKLELESKGGGSHPLLVPYDTL
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LFWGIFDSLSHKKYDPDLFRMALPCLSAIAGALPPDYLDTRISATLEKQISVDADGNF
                                                                                                                                                                LTRCAPLFAGTEHYTSLIDSTLQTIYRLSKGRSLTKAQRDTIEECLLAICNHLRPSML
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LFLDRVYGIKDQTFLLHLLEVGFLPDLRASASLDTVSLSTTEAALALNRYLCSAVLPL
                                                                                                                                                                                                                                                                                                                                                                                                                                       SRYNELMQALNMSAALTARKTREFRSPPQEQINMLLNFQLGENCPCPEEIREELYDFH
EDLLIHCGVPLEEEEEEEDTSWAGKLRALVYKIKGPPKPEKEQPTEEEERCPTTLKE
LISQTMIRWAQEDQIQDAELVRMMFSLLQRQYDSIGELLQALRKTYTISQASVSDTIN
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KLIGTLLVMGVFDDDDIRQILLLIDPSVFGEHSVETEDGAEKEEVTQVEEKAVEAGEK
AGKEAPVKGLLQTRLPESVKLQMCELLSYLCDCELQHRVEAIVAFGDIYVSKLQANQK
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DLMQFHYHTLRLYSAVCALGNSRVAYALCGHVDLSQLFYAIDNKYLPGLLRSGFYDLL
ISIHLANAKERKLMMKNEYIIPITSATRKIRLYPDESKRHGLPGVGLRTCLKPGFRFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HESVKRSNCYMVWGGDIVATSQRSSRSNVDLEIGCLVDLAMGMLSFSANGRELGTCYQ
VEPNTKVFPAVFLQPTSTALFQFELGKLKNAMPLSAAIFKSEEKNPVPQCPPRLDVQT
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MDSPPCLKVTHKTFGTQNSNASMIYCRLSMPVECHSSFSHSPCLDSEAFQKRKQMQEI
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EKNYNLQMSTETLKTLLALGCHIAHVNPAAEEDLKKVKLPKNYMMSNGYKPAPLDLSD
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IESPEALNLIAEGHIKSIISLLDKHGRNHKVLDVLCSLCLCNGVAVRANQNLICDNLL
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QPPEEEMQHEDKQNKLRSLKNRQNLFKEEGMLALVLNCIDRLNIYNSVAHFAGIAREE
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KRDIEGMGVPEIKYGDSVCFVQHIASGLWVTYKAQDAKTSRLGPLKRKVILHQEGHMD
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                                                                                                                                                                                                                                                                                                       EGERYLSFLRFAVFVNSESVEENASVVVKLLIRRPECFGPALRGEGGNGLLAAMQGAI
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/protein_id="CAA69029.1"
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db_xref="taxon:9667"

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TAKEKFRDREKAQDLFKFLQVNGIVVSRGVKDMELDASSMEKRFAYKFLKKILKYVDS AQEFIAHLEAIVSSGKTEKSPHDQEIKFFAKVLLPLVDQYFTNHHLYFLSSPLKPLSS

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RESULT 4
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CNS06JG2

Of Saccharomyces bayanus, sequence tagged site.

of Saccharomyces bayanus, sequence tagged site.

AL401544

AL401544.1 GI:12159319

STS.

Saccharomyces bayanus

Bukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycets

Saccharomycetales; Saccharomycetaceae; Saccharomycets;

Saccharomycetales; Saccharomycetaceae; Saccharomycets;

Souciet, J.L., Aigle, M., Artiguenave, F., Blandin, G.,

Bolotin-Fukuhara, M., Bon, E., Brottier, P., Casaregola, S.,

de-Montigny, J., Dujon, B., Durrens, P., Lepingle, A., Llorente, B.,

Malpertuy, A., Neuveglise, C., Ozier-Kalogeropoulos, O., Potier, S.,

Saurin, W., Tekaia, F., Toffano-Nioche, C., Wesolowski-Louvel, M.,

Wincker, P. and Weissenbach, J.

Genomic exploration of the hemiascomycetous yeasts: l. A set of

veast species for molecular evolution studies
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SGYASHKEKEMVASLFCKLAALVRHR ISLFGSDSTIMVSCLII ILAQTLDTRTVKKSGS
ELLVKAGLRAFFENAABDLEKTSENLKLGKFTHSRTQIKGVSONI NYTTVALLPILTSI
FEHVAQHQFGVDLLLGDVQISCYHILCSLYSLGTGKNIYVERQRPALGECLASLAAAI
PVAFLEPTLLMRYNPLSVENTKTPRERSILGHADTVEEWEQDIPOLEGLMKE ISDLAES
GARYTEMPHVIEVILPMLCNYLSYWMERGPEHLPSTGPWCTKVTSEHLSVILGULIK
IINNNLGIDEASWMKRTAVYAQPIISKARPDLLKSHFIPTLEKLKKKAVKTVQEEEQL
KADTKGDTQEAELLILDEFAVLCRDLYAFYPMLIRVDNNNSNMEKSPSADSDQLFRM
VAEVFILWCKSHNFKRESONFVIQNEN INLAFTLTGDSKSKMSKSGQDQERKKTKRRG
DLYSIQTSLIVAALKKMLPIGLNNCTPGDDELISLAKSRYSTRDTDEEYCBLLKK
INNNLGIDEASWMKRTAVYAQPIISKARPDLLKSHFIPTLEKLKKKAVKTVQEEEQL
KADTKGDTQEAELLILDEFAVLCRDLYAFYPMLIRVYDNNNSNMEKSPSADSDQLFRM
VAEVFILWCKSHNFKRESONFVIQNEN INLAFTLTGDSKSKMSKSGQDQERKKTKRRG
DLYSIQTSLIVAALKKMLPIGLNNCTPGDDELISLAKSRYSTRDTDEEYCBLLKK
INNNLGIDEASWMKRTAVYAQPIISKARPDLLKSTFORVELHENULK
LQEKSDDPAVKWQLNLYKOVLKSEEPFNAEKTIERVQRISAAVFHLEQVEQPLRSKKA
VWHKLLSKQRKRAVVACFRMAPLYNLPRHKINNFFLSTEFQRVWLEKVNEKTQYDRLIP
ILMKSPKVEEEEEEEEMEKKORTLYQQARLHERGAAEMYLQMISASGEMSPMVV
ETLKLGIAILNGGNAGVQOKMLDYLKEKKDAGFFOSLSGLMQSCSVLDLNAFERQNKA
EGLGMVTEEGTREKVLONDEFTRDLFRFLOLLCEGHNSSDFORVLETTINVII
STVDYLLTLQESISDFYWYYSGKDIIDESGQHNESKALAVTKQIFNSCTSTSKGEMSPMVV
EFOXHLMPANDSRLKCLLDPAGSVLNYFEPYLGRIEHMSDFQNFLRTQMGNTTTINVII
STVDYLLTLOESISDFYWYYSGKDIIDESGQHNESKALAVTKQIFNSCTULAFERQNKLLE
POVYLESKROFT TOVVNEGGSQEKMELEVNFCEDTIIEMGAKIERVYEEISESSRTQWEK
POVNESKROFT TOVVNEGGSQEKMELEVNFCEDTIIEMGAKIERVYEEISESSRTQWEK
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POVRESKROFT TOVVNEGGSQEKMELEVNFCEDTIIEMTVFGELOADIMSDLEGFHPKKE
EDGEKEDTAKEEEQAEALVALVTKKKRARAGAAEMKAAHEAEKVEREDLY
KRYWKELSTGAABALLDIAMGFKTLRTILSSTVRKKRAQAAEMKAAHEAEKVEREDLETYOTKLLHY
LARNEYNLLESTOMAPALRALAIHTVISLVCVVGYYCLKVPTDNSFDLAMYTTMSYLLHY
NNEFERAAHLLDIAMGFKTLRTILSSTVRKKHAQAAEMKAAHEAEKLEFDGLY
TEQOSEDDILEDNKGDEAGSLSSIDMKYHIMKLGVVFTDNSFDLAMYTTMSYLLFDLGLIK
NALDFFRAFLEDLGLIKGFDLATHATVTSLSVCVYFTDNSTALAWYTTMSYLLFDLGLIK
NALDFFRAFLEDLGLIKGFDLYTHOKYGLAVVTYNDRYDLYVARNEFTEKE
YNKSEDDDEPDMCDMKDATALAILTILSETDGLYTHOKYGLA
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FFFVIVILLAIIQGLIIDAFGELRDQQEQVREDMETKCFICGIGNDYFDTTPHGFETH
TLQEHNLANYLFFLMYLINKDETEHTGQESYVWKMYQERCWDFFPAGDCFRKQYEDQL
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cDNA encoding human G p

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E15997.1 GI:5710680

JP 1998127289-A/1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Aigle, M. and Durrens, P. Genomic exploration of the hemiascomycetous saccharomyces bayanus var. uvarum FEBS Lett. 487 (1), 37-41 (2000)
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Direct Submission
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OS Homo sapiens (human)
PN JP 1998127289-A/1
PD 19-MAY-1998
PF 29-OCT-1996 JP 1996286823
PI HINUMA KUNIJI, FUKUZUMI MASASHI, KAWAMATA YUJI PC
C12N15/09,C07H21/04,C07K14/705,C07K16/28,C12N1/21,C12P21/02,
C12P21/08,
PC C12Q1/68,G01N33/53,G01N33/566//A61K38/00,A61K39/395,
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                                                                                                                                                  a,K., Fukuzumi,M. and Kawamata,Y.
PROTEIN CONJUGATE TYPE RECEPTOR PROTEIN AND ITS DNA
t: JP 1998127289-A 1 19-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  TTGGATCGG 122
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Neuveglise,C
                                                                                                                                         CHEM IND
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/clone="AS0AA028E04"
/clone_lib="AS0AA"
/note="end : T7"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   complement(<18. .>422)
/note="similar to Saccharomyces
/note="similar to Saccharomyces
FPS1; glycerol channel protein
/evidence=not_experimental
213 c 243 g 328 t
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Location/Qualifiers
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                                                                                                                                                                                                                         Metazoa; Chordata; Craniata; Vertebrata; Eutheria; Primates; Catarrhini; Hominidae;
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. 1.8e+03;
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receptor pr
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                                                                                                                                                                                                                                                 Euteleostomi;
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DEFINITION
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E15998/c
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                                                                                                                                      OS Homo sapiens (human)
PN JP 1998127289-A/2
PD 19-MAY-1998
PF 29-OCT-1996 JP 1996286823
PI HINUMA KUNIJI, FUKUZUMI MASASHI, KAWAMATA YUJI PC
C12N15/09,C07H21/04,C07K14/705,C07K16/28,C12N1/21,C12P21/02, PC
C12P21/08,
PC C12Q1/68,G01N33/53,G01N33/566//A61K38/00,A61K39/395,
A61K39/395, PC A61K48/00,
PC (C12N1/21,C12R1:19),(C12P21/02,C12R1:19);
CC strandedness: Double;
CC hypothetical: No;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A61K39/395, P
PC (C12N1/2
CC stranded
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CC anti-sen
FH Key
FH Key
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                                                                                                                                                                                                                                                                       NEW G PROTEIN CONJUGATE TYPE RECEPTOR Patent: JP 1998127289-A 2 19-MAY-1998;
TAKEDA CHEM IND LTD
OS Homo cart
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                                                                                      Eukaryota; Metazoa; Chordata; Craniata; V. Mammalia; Eutheria; Primates; Catarrhini; 1 (bases 1 to 1626)
Hinuma, K., Fukuzumi, M. and Kawamata, Y.
                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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E15998.1 GI:5710681  
JP 1998127289-A/2.
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Location/Qualifiers
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Location/Qualifiers
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/db_xref="taxon:9606"
473 c 425 g 29
∕organism="Homo sapiens"
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/clone='pHEBF2'
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/tissue_type='brain
/clone='pHEBF2'
                                           /product='human
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Prpv 69/3
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Richards, J.G.

A new family of orphan G protein-coupled receptors

expressed in the brain

FEBS Lett. 424 (3), 193-196 (1998)
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AX01842
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larity 100.0%;
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/clone="lambda
52. .1497
                                                                                                                                                                                                             /translation="MRWLWPLAVSLAVILAVGLSRVSGAPLHLGRHRAETQEQQSRS KRGTEDEEAKGVQQYVPEEWAEYPRPIHPAGLQPTKPLVATSPNPDKDGGTPDSGQEL RGNLTGAPGQRLQIQNPLYPVTESSYSAYAIMLLALVVFAVGIVGNLSVMCIVWHSYYLKSAWNSILASLALWDFLVLFFCLPIVIFNEITKQRLLGDVSCRAVPFMEVSSLGVTTFSLCALGIDRFHVATSTLPKVRPIERCQSILAKLAVIWVGSMTLAVPELLLWQLAQEPAPTMGTLDSCIMKPSASLPESLYSLVMTYQNARMWWYFGCYFCLPILFTVTCQLVTWRVRGPPGRKSECRASKHEQCESQLNSTVVGLTVVYAFCTLPENVCNIVVAYLSTELTRQTLDLLGLINQFSTFFKGAITPVLLLCICRPLGQAFLDCCCCCCEECGGASEASAANGSDNKLKTEVSSSIYFHKPRESPPLLPLGTPC"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="lambda GT10"
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E 1 (bases 1 to 10585)

S Stephens, R.S., Kalman, S., Lammel, C.J., Fan, J., Marathe, R., Aravind, L., Mitchell, W.P., Olinger, L., Tatusov, R.L., Zhao, Q., Koonin, E.V. and Davis, R.W.

Genome sequence of an obligate intracellular pathogen of humans Chlamydia trachomatis

Science 282 (5389), 754-759 (1998)

99000809

9784136

E 2 (bases 1 to 10585)

S Kalman, S., Mitchell, W., Marathe, R., Lammel, C., Fan, J., Hyman, R. Olinger, L., Grimwood, J., Davis, R.W. and Stephens, R.S.

Comparative genomes of Chlamydia pneumoniae and C. trachomatis

Nat. Genet. 21 (4), 385-389 (1999)

99206606

D 10192388

E 3 (bases 1 to 10585)

S Stephens, R.S., Kalman, S., Lammel, C.J., Fan, J., Marathe, R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         157
                                                                                                                                                                                                                                                                                                                                                         AE001342
Chlamydia trachomatis s
AE001342 AE001273
AE001342.1 GI:3329166
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AX018426
AX018426.
                                                                                                                                                                                                                                                                                          Chlamydia
Chlamydia
Bacteria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Mammalia; Eutheria; 1 (bases 1 to 2411)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /codon_start=1
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SDNKLKTEVSSSIYFHKPRESPPLLPLGTPC"

1463. . 2411

37 a 775 c 659 g 580 t
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/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                             trachomatis
Chlamydiales; Chlamydiaceae;
1 to 10585;
                                                                                                                                                                                                                                                                                                                            trachomatis.
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Primates;
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section 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14; DB
Pred. No. 1.7
0; Mismatches
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Catarrhini;
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1.7e+03;
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of 87
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Tatusov, R.L., Zhao, Q.
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the complete
                                                                                                                                                                                                                pathogen of humans
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                                                                                                                  Hyman,R.W
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FEATURES
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10585
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Koonin,
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Berkeley, CA 9
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CA 94720-7360,
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, USA
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/codon_;
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                                                                                                                     REFERENCE
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                                                                                                                                                                                                                                                                                                                                                                                                             SOURCE
  FEATURES
                                                                      AUTHORS
TITLE
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                                                                                                                                                                                                                                                                     TITLE
                                                 JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM
                                                                                                                                                                                            JOURNAL
                                                                                                                                                                                                                                                                                            AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   00
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gene

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gene; tRN
tRNA-His
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AJ303116

AJ303116.1 GI:17932764

128 ribosomal RNA; 128 rRNA genome.
                                                       Direct Sub
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene; tR
tRNA-Ser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RNA-Leu(
transfer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           transfer
RNA-Cys;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ITS1;
                                                                                                                                                                                                                                                                                                                                                                                         Australian echiqna.
Mitochondrion Tachyglossus aculeatus
Enkaryota; Metazoa; Chordata; Crania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              dehydrog
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene;
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                          Univers
                                                                                                           Janke, A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; tRNA-Cys gene; tRNA-Gln g
-His gene; tRNA-Ile gene; t
; tRNA-Lys gene; tRNA-Met g
-Ser (AGY) gene; tRNA-Ser (
; tRNA-Tyr gene; tRNA-Val g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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AFPEMHPSVDIVPFLKPVAHKTSRLSSIVLSITIVLLSLGILGFATFYLAFHTYDHVS
QQKEKIQSINIPKLIEMMKRESPEKVALILSYLDSAKAEELLNKLPEEMKSAVLKLRT
"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 dehydrogenase subunit 3; NADH dehydrogenase subunit 4; NADH rogenase subunit 4L; NADH dehydrogenase subunit 5; NADH rogenase subunit 4; NADH gene; NADH2 gene; NADH3 gene; NADH4 rogenase subunit 5; NADH rogenase subunit 4; NADH4 rogenase subunit 5; NADH4 rogenase subunit 5; NADH4 rogenase subunit 4; NADH4 rogenase subunit 4; NADH4 rogenase subunit 4; NADH4 rogenase subunit 4; NADH4 rogena; NADH4 gene; NADH4 rogenasier RNA-Ala; gene; Lransfer RNA-Glu; transfer RNA-Gly; transfer RNA-Ser(UCN); transfer RNA-Ser(UCN); transfer RNA-Ser(UCN); transfer RNA-Ser (UCN); transfer RNA-Ser (UCN) gene; transfer RNA-Gly gene; transfer RNA-Leu (UUR) transfer transfer; transfer tr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NADH4L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NADH dehydrogenase subunit 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UUR); tran
Submission

ed (21-DEC-2000) Janke A., Deity, Soelvegatan 29, S-22362

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              enase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1914 c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NA; 12S rRNA gene; 16S ribosomal ounit 6; ATP synthase subunit 8; co2 gene; cox3 gene; cytb gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2513
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Pred. No. 1.6e+03;
0; Mismatches 0;
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(UCN)
gene.
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                                                        Dept.
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NADH dehydrogenase subunit 2;
                              Lund,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 10585;
                                SWEDEN
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                                                        Genetics,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA; 16S rR atp6 gene; cytochrome
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                                                                                                                                                                                                                                                                                                        1: AUC"
3796,aa:Ile)
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,aa:Phe)
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                                      : AUA"
3925,aa:Met)
                                                                                                                                                                                        CAA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ALLPLTPMFITLG"
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/gene="tRNA-Asn"
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                                                              FFDPAGGGDPILYQHLFWFFGHPEVYILILPGFGIISHIVTYYSGKKEPFGYMGMVWAMMSIGFLGFIVWAHHMFTVGMDVDTRAYFTSATMIIAIPTGVKVFSWLATLHGGDIKWTPPMLWALGFIFLFTVGGLTGIVLANSSLDIILHDTYYVVAHFHYVLSMGAVFAIMGGFVHWFPLLSGFTLHTTWAKVHFTLMFVGVNLTFFPQHFLGLAGMPRRYSDYPDAYTLWNAISSLGSFISLTAVMVMIFMVWEAFASKREVLTVELTSTNIEWLHGCPPPYHTFEEP
           complement(6937.
/gene="tRNA-Ser
                                              VYIKI"
       /gene=
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           . .6997)
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rce 155739 /organism="Arabido /strain="Columbia" /db_xref="taxon:37 /chromosome="3"	test information on annota kazusa.or.jp/kaos/cgi-bin similarity to proteins in or 'note' qualifiers. Gen milarity are described as re programs used to prediccs Group, Oak Ridge Nation pbio.ornl.gov/Grail-l.3/), hris Burge, MIT, http://CCS.M. Hebsgaard, et al., CBttp://www.cbs.dtu.dk/serviictor (Volker Brendel, Stamlinl.zool.iastate.edu/cgiding tRNAs are predicted b, Washington University Scome.wustl.edu/eddy/tRNAscance may not be the entire cause we remove overlaps b ne is MVAll and the 3' cloocation/Oualifiers	ko,T. Ka ctural an ctural an ence feat BAC clone Res. 7 (3 3099 bases 1 t ko,T. Ka ko,T. Ka itted (28 itted (28 itted (28 itted (28	AP00038 AP00038 AP00038 AP00038 Arabido Clone:M Arabido Clone:M Arabido	93.3%; Score 100.0%; Pred ative 0; Mi: CGGC 15 []] CGGC 9660	lement(6937. e="tRNA-Ser (duct="tRNA-Ser (duct="tRNA-Ser (pos:6)
/codon_start=1 /evidence=not_experimental /product="Mutator-like transposase" CDS 3323833464 /note="contains similarity to En/Spm-like transposon	xperimental B01421.1" 994246" 994246" STKKLFFPKRLFQP FAEMDLAFSSHIVH SPNVKRKKKEKWWI LRKQYGLFGFVQSI MIADGDKVHFLSCG QVEQGHQGRIDHVL EELEKVEYRGDEG EIPKQGNEEMEVEE NGTAPTEKESQPQK SQPQKEETTEVPRE DLRVKKWAKNKLIR KLLRAHPEVKYLVE (2376626453)) MSJ3.7"	568113/23,1380816049,1612/15233699717124,1722717382) 213.1 ty to unknown protein" xperimental 065,2014420368,2046720550,20807 180921955,2203622182,2226322353 262422794,2289422941) 853.1 ty to unknown protein"	<pre>/note="gene_id:MSJ3.3 unknown protein" /pseudo /codon_start=1 /evidence=not_experimental 8666. 9013 /note="gb AAF24530.1 gene_id:MSJ3.4 similar to unknown protein" /pseudo /codon_start=1 /evidence=not_experimental join(1342713562,1360513769,14564)</pre>	/db_xrer="G1:1194245" /translation="MDPNKAEISRLEVLPQDLLGEIVAKIGAKYAEDYHNCILSCKEL /translation="MDPNKAEISRLEVLPQDLLGEIVAKIGAKYAEDYHNCILSCKEL /complement(join(24204168,43285023,51385459)) /note="contains similarity to Arabidopsis thaliana retrotransposon Athila gene_id:MSJ3.2" /pseudo /codon_start=1 /evidence=not_experimental 67848561	sui F 1345. 810.1 0wn p

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ELAANERLSELPCRHYYHKECISNWLSNRNTCPLCRHNVELPNHG"
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/note="contains similarity to En/Spm-like transposon
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THSDPRETVLFNRFTPFDYIIEDTAHTENLVDLVGALVDVGCLRNITEDGWEMDGFRL
TFEIMDKYKERLKCEAFGELAVDLHRKFRSFRNDSIVIDNIFVALTWWKIFRYHTGLK
HVKFQNYAGISDVTVDPNIEEVQELKMVLE"
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RDNMRALKRDGRVV"
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                                                                                           * NOTE: This record contains 85 individual

* sequencing reads that have not been assembled into

* contigs. Runs of N are used to separate the reads

* and the order in which they appear is completely

* arbitrary. Low-pass sequence sampling is useful for

* identifying clones that may be gene-rich and allows

* overlap relationships among clones to be deduced.

* However, it should not be assumed that this clone

* will be sequenced to completion. In the event that

* the record is updated, the accession number will
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cambridgeshire, CB10 1SA, UK. E-mail enquiries:
humquery@sanger.ac.uk Clone requests: clonerequest@sanger.ac.uk
On Mar 1, 2002 this sequence version replaced gi:16973133.
During sequence assembly data is compared from overlapping clones.
Where differences are found these are annotated as variations
together with a note of the overlapping clone name. Note that the
variation annotation may not be found in the sequence submission
corresponding to the overlapping clone, as we submit sequences with
only a small overlap as described above.
This sequence was finished as follows unless otherwise noted: all
regions were either double-stranded or sequenced with an alternate
chemistry or covered by high quality data (i.e., phred quality >=
30); an attempt was made to resolve all sequencing problems, such
as compressions and repeats; all regions were covered by at least
one plasmid subclone or more than one M13 subclone; and the
assembly was confirmed by restriction digest. The following
abbreviations are used to associate primary accession numbers given
in the feature table with their source databases: Em:, EMBL; Sw:,
SWISSPROT; Tr:, TREMBL; Wp:, WORMPEP; Information on the WORMPEP
                                                                                                                   6184
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//www.chori.org/bacpac/home.htm
R: pBACe3.6
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  69928
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* identifying clones the
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nitted (23-NOV-2001) Whitehead Institute/MIT Center for Genome nitted (23-NOV-2001) Whitehead Institute/MIT Center for Genome earch, 320 Charles Street, Cambridge, MA 02141, USA repeats were identified using RepeatMasker:
repeats were identified using RepeatMasker:
t, A.F.A. & Green, P. (1996-1997)
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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40957: gap of 100 bp 41823
41822: gap of 100 bp in 42586
42685: gap of 100 bp 42586
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44311: contig of 750 bp in 4312
44411: gap of 100 bp 45261: gap of 100 bp 45262
46121: gap of 100 bp 46871
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47702: contig of 730 bp in 48543
48642: gap of 100 bp 48643
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31462: contig
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32432: gap of
33269: gap of
34111: gap of
34964: gap of
35718: contig
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1 (bases 1 to 73509)

Wallis, J.

Direct Submission
Submitted (16-OCT-2001) Sanger Centre, Hinxton, Cambridgeshire,
CB10 1SA, UK. E-mail enquiries: humquery@sanger.ac.uk Clone
requests: clonerequest@sanger.ac.uk
On Oct 21, 2001 this sequence version replaced gi:16214503.
                                                             ACCTCTTGGATCGG
                                                                                                                                                                                                                                                                               NOTE: This is a 'working draft' sequence This sequence will be replaced by the finished sequence as soon as it the accession number will be preserved.

Location/Qualifiers
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/db_xref="taxon:9606"
/chromosome="11"
/map="p12-14.1"
/clone="RP5-1002E13"
/clone_lib="RPCI-5"
1. 73509
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0; Mismatches 0;
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| SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1982.DAT:*
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392.790 million cell updates/sec
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Compugen
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Abl variable heavy G-protein coupled Human G protein-co DNA encoding human Human EST-derived G-protein coupled Human mucilage cod cDNA encoding a G-Human brain G-prot
                                                                                        Description
  RESULT 1
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DE Ab1
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                                                                                                                                    Homo
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                     Adams CW,
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                                       (GETH ) GENENTE
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ABK64653

ABL22291

AAH07127

AAF21557

AAF21557

AAF21557

AAAF21557

AAAA81272

AAAA81272

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RESULT 2
AAV27174/c
ID AAV27174
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Thrombocytopenia -associated bone marrow hypoplasia (e.g. aplastic anemia following chemotherapy or bone marrow transplant) may be effectively treated with the antibody compounds as well as disorders such as (HIV-induced and non HIV-induced), chronic idiopathic thrombocytopenia (congenital thrombocytopenia, thrombotic thrombocytopenia and congenital thrombocytopenia, thrombotic thrombocytopenia and myelodysplasia. They can also be used in e.g. myelotoxic chemotherapy for autologous or allogeneic bone marrow transplant, myelodysplasia, idiopathic aplastic anemia, congenital thrombocytopenia, and immune thrombocytopenia. The antibodies which bind to the MuSK receptor can be used for improving neuromuscular function in a patient, e.g. in muscular dystrophy. The products can also be used for detection and diagnosis. The sequences have a longer half-life than the natural ligand for the TPO-R. CDR2, and CDR3 regions of variable heavy (VH) chains and variable light (VL) chains of antibodies Abl to Ab6.
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Best Local S
Matches 15
            WPI; 19
P-PSDB;
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                                                  Fukusumi S,
                                                                                                      29-OCT-1996;
                                                                                                                              27-OCT-1997;
                                                                                                                                                          03-JUN-1998
                                                                                                                                                                                                                                       G-protein coupled receptor; short form; human; ds.
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DB; AAW55029.
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hematological disorders
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                                                                                 The invention re of endogenous hu AAY90677 and AAY and AAY
                   mutation in a portion of the protein comprising intracellular I (IC3) and transmembrane domain 6 (TM6). A non-endogenous amino is substituted for an endogenous residue in IC3 at a position I
  is substituted acids N-termina
                                                                                                                                                                                                                                     receptors, agents -
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P-PSDB; AAY9063
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  or gene therapy for diseases caused by a deficiency of the DNA can also be used to detect abnormalities in the gene receptor. The protein or fragment can be used to determine eptor ligands in vivo. The antibody can be used in assays to
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                                                                                elates to constitutively active, non-endogenous versions uman orphan G protein-coupled receptors (GPCRs, AAY90643~Y90683-Y90687), and to DNA encoding them (AAA30709-A307430779). The mutant proteins of the invention contain a
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ID AAA30731 s

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DT 21-AUG-201

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DE DNA encod

XX

CO protein

KW intracelli

KW agonist;

XX

OS Homo sapid

OS Synthetic

XX

PN WO2000221

XX

PD 20-APR-20

XX

PF 12-OCT-19

XX

PF 12-OCT-19

XX

DR WPI; 2000

DR P-PSDB; A

XX

DR WPI; 2000

DR P-PSDB; A

XX

PT agents

PT receptors

PT agents

PS Example 2

XX

CC The inven

CC AAY90677

CC and AAA30

CC mutation

CC mutation

CC TOTOLOGY

CC mutation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C X-(AA)15-Pro. The endogenous amino acid is selected from Lys, His, Arg or Ala, and is preferably Lys. When the endogenous residue at this C position is Lys, this residue is replaced by His, Arg or preferably Ala. The 15 amino acid stretch between the substituted amino acid and the Pro may be endogenous, non-endogenous, or a mixture of endogenous and cidentifying antagonists, agonists and partial agonists for use as C pharmaceutical agents. The mutant proteins are also useful in research cidentings for elucidating the roles of the receptors in normal and C diseased conditions. Antagonists for a particular GPCR are useful for treating diseases and disorders associated with that receptor. Because the novel mutant GPCRs are constitutively active, they can be used C directly for screening of compounds without the need for endogenous cligands. The present sequence represents cDNA encoding a human wild-type can be detected to site-directed mutagenesis (SDM) to generate DNA encoding corresponding mutant of the invention.
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Best Local
Matches 1
                                                                                                                                                                                                                                                                                                                                                           12-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      G protein-coupled receptor; GPCR; cointracellular loop 3; transmembrane
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  The invention relates to constitutively active, non-endogenous versions of endogenous human orphan G protein-coupled receptors (GPCRs, AAY90643-AAY90677 and AAY90683-Y90687), and to DNA encoding them (AAA30709-A30743 and AAA30775-A30779). The mutant proteins of the invention contain a mutation in a portion of the protein comprising intracellular loop 3 (IC3) and transmembrane domain 6 (TM6). A non-endogenous amino acid, X,
                                                                                                                                                                                                                                                                                                                                                                                                                          WO200022129-A1
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                                                                                                                                                       receptors,
                                                                                                                                                                              Non-endogenous
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ful for identifying agonists for
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Pred. No. 2.2e+02;
Mismatches 0;
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ne domain 6; drug screening;
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RESULT 5
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ID AAH98401:
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AC AAH98401:
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AC AAH98401:
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DT 12-OCT-20
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Human EST
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Human; sh
KW Human; sh
KW diagnosti
KW biodivers
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OS Homo sapi
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PN W02001544
XX
PD 02-AUG-20
XX
PF 25-JAN-20
PR 15-SEP-20
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PJ Cao Y, D
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PI Tang YT,
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PT antibodie
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tomato; monkey;
diagnostics; for
biodiversity; ge
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 l of an endogenous proline in TM6 to form a sequence he endogenous amino acid is selected from Lys, His, preferably Lys. When the endogenous residue at this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ig; cow; fruit fly; yeast; hamster; macaque; h
  dog; sea urchin; expressed sequence tag; EST;
rensic test; gene mapping; genetic disorder;
ene therapy; nutrition; ss.
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2.2e+02;
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25-JAN-2000; 17-JUL-2000; 03-AUG-2000; 15-SEP-2000; 25-JAN-2001; HYSEQ Drmanac Liu 2000US-0491404. 2000US-0617746. 2000US-0631451. 2000US-0663870. 2001WO-US02687. a RΑ, Zhou 2hang ρ, Qian XB, J, Werhn Werhman T; Wang Z, Chen **χ**,

The present invention provides the protein and coding sequences of novel proteins from a variety of organisms, including human, dog, cat, horse, cow, pig, hamster, monkey, macaque, yeast, bacteria, fruit fly, sea

from a vai

antibodies

and

research use

polypeptide

for treatment of diseases,

diagnostics,

Page

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68-369; 1275pp; English.

2001-47616 DB; AAM2374

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RESULT 6
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AAV27175
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                                                                                                                                                                         invention. The protein or cells expressing the DNA encoding it can be used to screen for agonists or antagonists of the receptor, which can used as drugs for treating various diseases (none disclosed). The DNA also be used for practice drug design based on comparisons with structurally analogous ligands and receptors. DNA encoding the protein can be used for gene therapy for diseases caused by a deficiency of the protein can be used for gene therapy for diseases caused by a deficiency of the protein can be used for gene therapy for diseases caused by a deficiency of the protein can be used for gene therapy for diseases caused by a deficiency of the protein can be used for gene therapy for diseases caused by a deficiency of the protein can be used for gene therapy for diseases caused by a deficiency of the protein can be used for gene therapy for diseases caused by a deficiency of the protein can be used for gene therapy for diseases caused by a deficiency of the protein can be used for gene therapy for diseases caused by a deficiency of the protein can be used for gene therapy for diseases caused by a deficiency of the protein can be used for gene therapy for diseases caused by a deficiency of the protein can be used for gene therapy for diseases.
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                                                                                                                                   levels
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                                                                                                Sequence
                                                                                                                                     receptor. The DNA can also encoding the receptor. The levels of receptor ligands
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                                                                                                                                                                                                                                                                                   Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EP845529-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -protein
                                                                                                                                                                                                                                                                                                          human G-p
, ligands,
                                                                                                                                                                                                                                                        sequence
                                                                                                                                                                                                                                                                                                                                                1998-288746/26
)B; AAW55030.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  form;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  166
                                                                                                                                                                                                                                                                                   ٠.
ص
                                                                                                                                                                                                                                                                                                                                                                                                             ) TAKEDA CHEM
                                               Similarity 100.0%;
14; Conservative
                                                                                                                           of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity 100 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Š
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CCTCTTGGATCGGC
                                                                                                                                                                                                                                                                                 Page
CCTCTTGGATCGGC
                        CCTCTTGGATCGGC
                                                                                                   1626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                         G-protein coupled receptor nds, antibodies, etc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                coupled
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human;
                                                                                                                           protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first
                                                                                                                                                                                                                                                                                                                                                                                      Hinuma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP;
                                                                                                 B₽;
                                                                                                                                                                                                                                                        encodes a
                                                                                                                                                                                                                                                                                 50-51;
                                                                                                                                                                                                                                                                                                                                                                                                                                      96JP-0286823.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                97EP-0308562
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      269
                                                                                                 272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       receptor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                              IND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                      Ś
                                                                                                                                                                                                                                                                                 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            93.3%;
100.0%;
                                                                                                 À;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ?
308
                        <u>1</u>5
                                                                                                                                                                                                                                                        human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15
                                                                                                                                                                                                                                                                                                                                                                                      Kawamata
                                                                                                 540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      526
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1626
                                                                                                                                      be used to detect abnormalities in the gene protein or fragment can be used to determine in vivo. The antibody can be used in assays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       long
                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;
                                                             Score 14;
Pred. No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14;
                                                                                                                                                                                                                                                        G-protein coupled
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ₽P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred.
                                             Misma
                                                                                                   484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                      Κ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       form,
                                      No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      G;
                                                                                                                                                                                                                                                                                                                      protein
                                           DB 19,
2.2e+02;
-~~ 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 22,
2.2e+02;
0;
                                                                                                   330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       n expressed sequence takes used in diagnostics, futations, to assess. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              abnormality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence.
                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                     and corresponding
                                                                                                                                                                                                                                                        receptor
                                                                        Length 1626;
                                                                                                other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 1616;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence
                                                                                                                                                                                                                                                        O.f.
                                                                                                                                                                                                                                                        the
                                                0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tags
                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         დ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cDNA
                                                                                                                                                                                                                  can
                                                                                                                                       C
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RESULT 7
AAZ11093/c
ID AAZ11093;
XX
AC AAZ11093;
XX
DT 03-NOV-199
XX
DE Human muc:
XX
DE Human muc:
XX
COS Homo sapio
XX
FH Key
FT CDS
FT
mucilage, while ante
peptides) of the pro
Typical of very many
bacterial, fungal, p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mucilage; obesity; a
                                                                                                                                                                                                                         of the protein is used to diagnose these diseases, or susceptibility them. The mucilage protein is also used to screen for specific (ant)agonists, which are potential therapeutic agents; to raise antibodies (Ab); in protective vaccines; to isolate cognate receptors and in structure based drug design. The mucilage coding sequences are used to produce recombinant mucilage; as a source of primers and probabilities.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disease; hypotension infarction; asthma; a Detecting mutations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This sequence encodes the mucilage protein of the invention. Mucilage is a G-protein-coupled receptor. Mucilage, or the nucleic acid encoding it, are used to treat conditions requiring increased activity/expression of mucilage, while antagonists (or nucleic acid encoding them or competitive peptides) of the protein are used where the opposite effect is required.
                                                                     are used to : purification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New mucilage re
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-MAY-1998;
20-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            osteoporosis; neurological
Sequence
                                                                                                                                                for isolating reidentification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     virus-1 or HIV-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-527457/44.
P-PSDB; AAY32886.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Elshourbagy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SMIK)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9942485-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mucilage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SMITHKLINE BEECHAM CORP.
    1728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         anorex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         human
                                                                                                                   ide
                                                                               of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              man; G-protein-coupled receptor; infection; cancer; diabet
rexia; Parkinson's disease; hypotension; hypertension;
stroke; myocardial infarction; asthma; allergy; therapy;
disorder; dyskinesia; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           receptor polypeptide for treatment, prevention and e.g. infections and cancer % \left( 1\right) =\left( 1\right) +\left( 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                              related genes or detecting mutations; for chromosome n; and as source of antagonistic oligonucleotides. The Ab dentify or isolate mucilage-expressing cells, for affinity of mucilage and the control of mucilage.
BP; 284 A; 569 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     34; 50pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98US-0075468.
98US-0075307.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2); pain; cancer; diabonsion or hypertension; hma; allergy; neurolog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99WO-US03320.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ions in the
                                                                        mucilage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  many diseases that can be treated are infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vawter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
150..1595
/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein are usemany dir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /product= mucilage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protozoal,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cdna;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   zoal, particularly human immune deficiency cancer; diabetes; obesity; anorexia; Parkinson's hypertension; osteoporosis; stroke; myocardial rgy; neurological disorders; dyskinesia etc. he DNA and/or determining the presence or amount
                                                                        and therapeutically.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1728 BP
513 G; 362 T; 0 other;
                                                                                                                                                                                                                                                                                                                    receptors;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       competitive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (viral,
                                                                                                                                                                                                                                                                                                                                                                                                                                             5
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Query Match Best Local S Matches 14

l Simil 14; C

arity

93.3%;

Score 14; DB 20; L. bred, No. 2.2e+02;

Length 1728;

Indels

0,

Gaps

0;

Conservative

S

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15

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RESULT 8
AAZ36408/c
ID AAZ36408;
XX
AC AAZ36408;
XX
DT 22-FEB-200
XX
DE CDNA encount
XX
KW Human; G-1
KW pathologic
XX
OS Homo sapic
XX
ST CDS
FT CDS
FT CDS
                                                                                      В
멍
                    9
                                          Query Match
Best Local S
Matches 14
                                                                                                                                        The present sequence encodes a human G-protein coupled receptor, designated HG01. HG01 is highly expressed in the brain, and is thought to be related to the endothelin receptors. The HG01 cDNA sequence was isolated from a human foetal brain library. HG01 can be used to identify binding substances, antagonists and agonists, which may be useful pharmacologically to modulate HG01 activity. G-protein coupled receptors (GPCRs) are involved in many pathological states, and modulators of GPCR activity are therefore useful to treat such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mat_peptide
                                                                                                          modulators of GPCR activity are therefore useful to treat sconditions. HG01 (or fragments) are also useful to produce The polynucleotides can be used in gene therapy, especially conditions requiring increased HG01.
                                                                                                                                                                                                                                                                                                     P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; G-protein pathological sta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ36408;
                                                                                                                                                                                                                                              Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                   21-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                        17-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9960153-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sig_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-FEB-2000
                                                                                                                                                                                                                                                                               Novel receptor
                                                                                                                                                                                                                                                                                                                                                           (MERI ) MERCK
                                                                                                                                                                                                                                                                                                                                                                                                                            25-NOV-1999
                                                                                     Sequence
                                                                                                                                                                                                                                                                   drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     encoding
                                                                                                                                                                                                                                                                                                     2000-053303/04
)B; AAY53779.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               287
                                           14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CCTCTTGGATCGGC
                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
                                                                                                                                                                                                                                              Fig 1A;
                                                                                                                                                                                                                                                                    development
                                                                                     1850 BP;
CCTCTTGGATCGGC
                    CCTCTTGGATCGGC
                                          93.3%;
larity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       D
                                                                                                                                                                                                                                                                                                                                                           δ'n
                                                                                                                                                                                                                                                                       protein, useful
                                                                                                                                                                                                                                                                                                                                                                                 98US-0086294
                                                                                                                                                                                                                                                                                                                                                                                                        99WO-US10808
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 159. 1526
/*tag= c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      coupled
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       G-protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
84..1529
/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /transl_except=
84..158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /product= "G-protein coupled receptor HG01"
/transl_except= (pos: 186..188, aa: Ala)
                                                                                                                                                                                                                                              33pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         note≖ "specifically"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag=
                                                                                      305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cDNA;
                                                                                      >
                                                                                                                                                                                                                                              English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ۵
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 274
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    receptor; therapy; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1850
                                                                                      614 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     coupled
                                          0;
                                                     Score 14;
Pred. No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ЧB
                                                                                                                                                                                                                                                                             for identifying antagonists
                                   Pred. No. 2.3e+02;
Mismatches 0:
                                                                                     540
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                                                                                      other;
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                                          Gaps
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RESULT 9
AAZ00860/c
PT PT PT
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LT.
H.A.
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general current, (2) as p...

(11) in usual hybring in us
                                            drug addiction e obesity, anorexi G-protein couple
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genes. (I) is used (
for specific ligands
therapeutic agents)
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5'UTR
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                                                                                                                                                      (I) are used to oparticularly for diseases of the o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   analgesic, anti-e anti-bulimic and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim
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                                                                                                                                 or Parkinson'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New human G-pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-FEB-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human
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)B; AAY2596
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                                         incy states, (2) as probes and primers for diagnostic (II) in usual hybridization/amplification assays, (3) for nt expression of (I), (4) to isolate the corresponding genomic (5) to design ribozymes and antisense sequences for treating ssion of (I), (6) to produce transgenic animals for studying ophysiology of (I) and (7) to detect mutations in (I)-encoding is used (1) to raise specific antibodies (Ab), (2) to screen fic ligands, agonists and antagonists, (potentially useful as ic agents) and (3) to treat (I)-deficiency conditions. Abs to seed to detect/determine (I) in standard immunoassays, rly for diagnosis. (I) is associated with chronic and acute of the central and peripheral nervous systems (e.g. Alzheimer's son's diseases, brain tumours, pain, schizophrenia, epilepsy, ction etc.), diseases that include a nervous component (e.g. anorexia and bulimia). This sequence encodes the human brain coupled receptor described in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ed receptor; human; brain; anti-neurodegeneration; pain; lgesic; anti-epileptic; anti-addictive; anti-obesity; anti-bulimic; cerebroprotective; gene therapy; diagnostic; ase; central nervous system; peripheral nervous system; ease; Parkinson's disease; brain tumour; schizophrenia; addiction; obesity; anorexia; bulimia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ystem disorders, e.g. Alzheimer's cerebral protectant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tein coupled receptor from brain tissue,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             rd;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             describes a novel G-protein coupled receptor isolated n tissue which has anti-neurodegeneration, antitumour epileptic, anti-addictive, anti-obesity, anti-anorex d cerebroprotective activity. Nucleic acid (II) that its complement, is used (1) in gene therapy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   erbach
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cerebroprotective activity. Nuclear ts complement, is used (1) in gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cDNA to mRNA; 2411 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disease,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               anti-anorexic,
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Sequence

2411

ВP;

397

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775 C;

659 G;

580 T;

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other;

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RESULT 10
AAZ01425/c
ID AAZ01425
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Best Local S
Matches 14
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Best Local S
Matches 14
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28-NOV-1997;
17-DEC-1997;
                                                                                                                            The present sequence represents the complete genome of Chlamydia trachomatis. Open reading frames (ORFs) of the genome encode polypeptides AAY36754-Y37949. The polypeptides can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal uretritis, epidymitis, cervicitis, salpingitis, perihepatitis, bartholinitis; pneumopathy in breast feeding infants; and venereal lymphogranulomatosis. The polypeptides of the invention may be of use in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vaccine; eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ01425;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Complete
                                                                                                                                                                                                                                                                         Claim 1;
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                                                                                           Sequence
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                                                                                                                   treating
683576
                                                                                                                                                                                                                                                                                                                                                                   ) GENSET
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                                               14;
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CCTCTTGGATCGGC
                                                         Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
                      ACCTCTTGGATCGG 14
                                                                                                                     these diseases.
 ACCTCTTGGATCGG
                                                                                             1038602
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                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
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97FR-0015041.
97FR-0016034.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence of
                                                                                                                                                                                                                                                                                               of Chlamydia
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683563
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Pred. No.
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                                                         Score 14;
Pred. No
                                                                                                                                                                                                                                                                                                trachomatis
                                                                                             214645
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                                             red. No. 3.5e+02;
Mismatches 0;
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2.3e+02;
0;
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                                                                   Length 1038602;
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                                             (Gaps
                                                                                            other;
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RESULT 11 AAS34099/c

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Human;
                Human cDNA encc
                                 17-DEC-2001
                                                 AAS34099;
                                                                  AAS34099
foetal
                                                                  standard;
                                (first
                oding
issue
                                                                  cDNA;
                a novel
                                                                  329
                foetal antigen,
                                                                   дВ
                SEQ ID No
                  623.
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Human; foetal tissue antigen; ss; antiinflammatory; neuroprotective; immunomodulator; cardiovascular; cytostatic; nephrothropic; cardiovascular; autoimmune disease; rheumatoid arthritis; hyperproliferative disorder; breast neoplasm; cancer; cardiovascular disorder; cardiac arrest; cerebrovascular disorder; cerebral ischaemia; angiogenesis; nervous system disorder; alzheimer's disease; infection; ocular disorder; corneal infection; wound healing; epithelial cell proliferation; food additive.

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AC AAS34099 standard; cDNA; 329 BP.

XX XX XX AAS34099;

XX 17-DEC-2001 (first entry)

XX 17-DEC-2001 (first entry)

XX 17-DEC-2001 (first entry)

XX Human: foetal tissue antigen: ss;

XX hyperproliferative disorder; braek

XX cardiovascular; autcimmune diseas

XX hyperproliferative disorder; braek

XX cardiovascular; autcimmune diseas

XX Wo200155312-A2.

XX I-JAN-2000; 2000US-0188664.

PR 14-JUL-2000; 2000US-0188664.

PR 14-JUL-2000; 2000US-0188664.

PR 12-JUL-2000; 2000US-0188664.

PR 12-JUL-2000; 2000US-0188874.

PR 11-JUL-2000; 2000US-01888974.

PR 11-JUL-2000; 2000US-02188990.

PR 11-AUG-2000; 2000US-02188990.

PR 11-AUG-2000; 2000US-02189874.

PR 11-AUG-2000; 2000US-0225213.

PR 11-AUG-2000; 2000US-0225266.

PR 11-AUG-2000; 2000US-0225275.

PR 11-AUG-2000; 2000US-0225275.

PR 11-AUG-2000; 2000US-0225297.

PR 11-AUG-2000; 2000US-0225299.

PR 12-AUG-2000; 2000US-022599.

PR 12-AUG-2000; 2000US-022599.

PR
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RESULT 12
AAQ66413/c
ID AAQ66413
XX
AC AAQ66413
XX
DT 26-JAN-1
XX
DE alcA pro
                                                                                                                                                                                                                                                                                     The invention relates to novel nucleic acids encoding novel human foetal cantigens. The nucleic acids and proteins are used to prevent, treat (e.g. to provent therapy) or ameliorate a medical condition in e.g. humans, mice, crabbits, goats, horses, cats, dogs, chickens or sheep. They care also used in diagnosing a pathological condition or susceptibility to a pathological condition. The antibodies to the antigens can also be used in alleviating symptoms associated with the disorders and in conclude autoimmune diseases e.g. radioimmuneassays or enzyme linked include autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cardiac arrest, viruses and fungi and ocular disorders e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to grevent skin aging due to sunburn, to maintain organs before as a food additive or preservative to increase or decrease storage capabilities, fat content, lipid, protein, carbohydrate, vitamins, camples of diseases and other nutritional components. Numerous candiscent are given in the specification. The present sequence.
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Best Loc
Matches
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17-NOV-2000;

01-DEC-2000;

01-DEC-2000;

05-DEC-2000;

05-DEC-2000;

06-DEC-2000;

08-DEC-2000;

08-DEC-2000;

08-DEC-2000;

08-DEC-2000;

08-DEC-2000;

11-DEC-2000;

11-DEC-2000;
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preventing or pr
musculoskeletal,
respiratory syst
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P-PSDB; AAU2127
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14; Conservative
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ACCTC:
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2000US-0249300.
2000US-0250160.
2000US-0251030.
2000US-0251988.
2000US-0251988.
2000US-0251479.
2000US-0251868.
2000US-0251869.
2000US-0251989.
2000US-0251990.
2000US-0259678.
                                                                                             nd
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ides and polypeptides for diagnosing, treating, rognosing e.g. diseases or disorders of the nervous, excretory, gastrointestinal, reproductive, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2/53.
                                  first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sh SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENOME SCI INC
                                                                                              ard; DNA;
                                                                                                                                                                         TTGGATTGGC 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tems
                                                                                                                                                                                                      TGGATCGGC 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          No 623; 642pp; English.
                                                                                                                                                                                                                                                   93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ruben SM;
                                                                                                 360 BP.
                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                   Pred.
                                                                                                                                                                                                                                                    Score 13.4; DB 22
Pred. No. 4.4e+02;
                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                    22;
                                                                                                                                                                                                                                      <del>---</del>
                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                         Indels
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                                                                                                                                                                                                                                         Gaps
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08-SEP-2000;
18-SEP-2000;
11-SEP-2000;
14-SEP-2000;
14-SEP-2000;
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14-SEP-2000;
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14-SEP-2000;
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14-SEP-2000;
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21-SEP-2000;
25-SEP-2000;
25-SEP-2000;
25-SEP-2000;
27-SEP-2000;
29-SEP-2000;
29-SEP-2000;
29-SEP-2000;
08-NOV-2000;
08-NOV-2000;
08-NOV-2000;
08-NOV-2000;
17-NOV-2000;

2000US-0231414.
2000US-0232081.
2000US-0232308.
2000US-0232399.
2000US-0232400.
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2000US-0233064.
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2000US-0249214.
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2000US-0249264.
2000US-0249264.
2000US-0249265.

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RESULT 13
AAS03486
ID AAS03
XX
AC AAS03
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DT 29-AU
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DE DNA e
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KW Antik
KW heart
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PN WO20(
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PP 11-O(
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DR WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bp
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Best Local S
Matches 14
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                                               WPI; 2001-282031
P-PSDB; AAU02586
                                                                                                                                                                                                                                                                                                                                                                                   12-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA constructs comprising a filamentous having a disrupted crea binding site - protein in the presence of glucose
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9413820-A
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                                                                                                                                                                                                Edwards
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA encoding anti-adipocyte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAS03486;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-JUN-1994.
                                                                                                                                                                                                                                                                                       (CAMB-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antibody; adip
heart disease;
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                                                                                                                                                                                                                                                                                       CAMBRIDGE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACCTCTTGGATCGGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               adipocyte;
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                                                                                                                                                                                                Main
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ВP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000WO-GB03900
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                                                                                                                                                                                                                                                                                                                                                                                   99US-0158812
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cyte; heavy chain; light chain; obes complementarity determining region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93WO-EP03553
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                                                                                                                                                                                                SH,
                                                                                                                                                                                                                                                                                       ANTIBODY TECHNOLOGY.
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Pred.
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ed. No. 4.4e
Mismatches
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Tateishi
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 Novel polynucleotides derived from Coryneform bacteria, for identic mutation point of a gene, measuring expression of a gene, analysing expression profile or pattern of a gene and identifying homologous
expression prof
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Senoh A, Ik

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ity 93.3%; Pred. No. 4.4e+02;
servative 0; Mismatches 1;
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Note: The sequence data for this patent did not form part of the priresectification, but was obtained in electronic format directly from the sequence is a nucleic acid describes and the prirect of the prire
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P-PSDB;
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AASO3401-AASO3535 represent anti-adipocyte monoclonal antibody heavy and light chain coding sequences of the invention. The antibodies can used in the treatment of obesity and obesity related diseases. The antibodies can be used to deliver drugs or pro-drugs directly to the feases of an obese patient or the antibody can be used as a therapeutic itself. Antibodies binding specifically to adipocytes can be used to activate the immune system to destroy the cells by complement mediated lysis. The antibodies may be labeled with a detectable label such as radiolabel, fluorescent or chemical group and used in methods of diagnosis in human subjects e.g. to determine the presence of adipocyte antigen on the surface of an adipocyte to detect or determine the
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REGISTRATION NUMBER: 23,031
REFERENCE/DOCKET NUMBER: TAK-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-407-0700
TELEFAX: 610-407-0700
TELEX: 846169
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
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Best Local
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           FILING DATE: 29-OCT-1996
ATTORNEY/AGENT INFORMATION:
NAME: Prestia, Paul F
                                                                                                             COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/959,381A
                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 286823/1996
FILING DATE: 29-0CT-1996
                                                                                                                                                                          ZIP: 19482
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
COMPUTER: IBM Compat
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COMPUTER
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STRANDEDNESS: single
TOPOLOGY: linear
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CITY: Valley
STATE: PA
COUNTRY: USA
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                                                                                     CLASSIFICATION:
                                                                                                     FILING DATE:
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MEDIUM TYPE: Diskett
COMPUTER: IBM Compat
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NUMBER:
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INFORMATION FOR
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MOLECULE TYPE:
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TELEPHONE: 6
                                                                                                 REFERENCE/DOCKET NUMBER: RP TELECOMMUNICATION INFORMATION: TELEPHONE: 864-233-6700
TELEFAX: 864-233-2284
FORMATION FOR SEQ ID, NO: 2:
                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/DE94/00022
FILING DATE: 12-JAN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P 43 01 017
                         MOLECULE
                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 39 base pairs
                                                                                                                                                                                          FILING DATE: ATTORNEY/AGENT
                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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        TYPE: nucleion STRANDEDNESS: TOPOLOGY: line OLECULE TYPE: DESCRIPTION:
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FILING DATE: 29
CLASSIFICATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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NUMBER: 26,
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16-JAN-1993
INFORMATION:
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N: 435
        other nucleic acid
/desc = "Oligonucleotide"
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RESULT 6
US-08-470-958-11/c
; Sequence 11, Application
; Patent No. 5674707
; GENERAL INFORMATION:
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                                                                                                                                                                                              ; DESCRIPTION: US-08-448-619-3
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Patent No
GENERAL
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Best Local
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/DE94/00022
FILING DATE: 12-JAN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P 43 01 017.2
FILING DATE: 16-JAN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:

CORRESPONDENCE ADDRESS:

ADDRESSEE: Hardaway Law F
                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: 864-233-6700
TELEFAX: 864-233-2284
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 29603-0107
COMPUTER READABLE FO
MEDIUM TYPE: Flo
COMPUTER: IBM PC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, 1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/448,619
FILING DATE: 29-SEP-1995
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LENGTH: 3> _
TYPE: nucleic actu
STRANDEDNESS: single
TOPOLOGY: linear
TOPOLOGY: linear
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STREET: P.
CITY: Gree
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14; Conservative
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VENTION: METHOD FOR THE OBTENTION OF NATIVE
VENTION: DOMAINS OF VIRAL MEMBRANE PROTEINS,
VENTION: AS VACCINE AGAINST HIV, AND THESE N
VENTION: MEMBRANE PROTEINS THEMSELVES
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SYSTEM: PC-DOS/MS-DOS
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93.3%;
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93.3%;
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= "Oligonucleotide"
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Federal Station
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Pred. No. 64;
0; Mismatches
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Pred. No. 64;
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GENERAL
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Best Local
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INFORMATION FOR SE
                                              STATE: V115-
COUNTRY: U.S.A.
COUNTRY: U.S.A.
ZIP: 22201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
COUNTYPEM: PC-DOS/MS-DOS
COUNTYPEM: PC-DOS/MS-DOS
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REFERENCE/DOCKE
TELECOMMUNICATION
TELEPHONE: (703)
TELEFAX: (703)
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ATTORNEY/AGENT IN
NAME: MITCHARD
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                         APPLICANT: Hintz, William E.
APPLICANT: Lagosky, Peter A.,
TITLE OF INVENTION: PRODUCTION OF HETEROLOGOUS PROTEINS
TITLE OF INVENTION: FILAMENTOUS FUNGI
             CURRENT APPLICATION DATA: APPLICATION NUMBER: US FILING DATE:
                                                                                                                                                                                               NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRE
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CITY: Arlington
STATE: Virginia
COUNTRY: U.S.A.
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TOPOLOGY: lir
OLECULE TYPE:
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TYPE: r
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
JRRENT APPLICATION DATA:
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CLASSIFICATION:
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AL INFORMATION:
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8TH FLOOR, 1100 NORTH GLEBE ROAD
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LAGOSKY, PETER A.
FENTION: PRODUCTION OF HETEROLOGOUS PROTEINS
FENTION: FILAMENTOUS FUNGI
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                                                                                                                                                                         Floor, 1100 No.
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EQ ID NO:
                                                                                                                                                                                                   ADDRESS:
                                                                                                                                                                                                                                                                                                          cation US/08321474
                                                    tentIn Release #1.0,
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BER: 29,009
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                            US/08/321,474
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Pred. No. 66;
0; Mismatches
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                                                     Version
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GENERAL INFORMATION:

APPLICANT: FLEISCHMAN, Robert D.

APPLICANT: WHITE, Owen R.

APPLICANT: FRASER, Claire M.

APPLICANT: VENTER, John C.

TITLE OF INVENTION: DNA SEQUENCES FOR STRAIN

TITLE OF INVENTION: TUBERCULOSIS

FILE REFERENCE: 24366-2007.00

CURRENT APPLICATION NUMBER: US/09/103,840A

CURRENT FILING DATE: 1998-06-24
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US-08-841-349-11/c
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US-09-103-840A-2
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REFERENCE/DOCKET NUMBER: 617-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 816-4000
TELEFAX: (703) 816-4100
TELEFAX: 200797 NIXN UR
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERS
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APPLICANT: MISHRA, LOPA
TITLE OF INVENTION: GENES CODING PROTEINS FOR EARLY
FILE REFERENCE: XX/PO4470US0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                             Sequence 2, Application US/09103840A Patent No. 6294328
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SEQ ID NO 11
LENGTH: 1177
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NUMBER
                                                                                                                                                                                                                                                                                                                           ORGANISM: Mus musculus FEATURE:
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APPLICATION NUMBER:
FILING DATE: 10-DEC-
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LENGTH: 360 base naing
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ATTORNEY/AGENT INFORMATION:
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ID NOS:
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; FEATURE:
; OTHER INFORMATION: C
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US-09-103-840A-2
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US-08-880-179-1
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APPLICANT: FLEISCHMAN,
APPLICANT: WHITE, Owen
APPLICANT: FRASER, Cla
                                                                                                                                                                                      Sequence 1,
Patent No. 6
GENERAL INE
                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE:
SEQ ID NO 1
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TITLE OF INVENTION: TUBERCULOSIS
FILE REFERENCE: 24366-20007.00
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE:
LITLE OF INVENTION:
LITLE OF INVENTION:
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESSEE:
STREFT
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TYPE: DNA
ORGANISM: Mycobacterium tuberculosis
                                                   APPLICANT: VOS,
TITLE OF INVENTI
TITLE OF INVENTI
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No. 6294328
                                                                                                                                                                                      INFORMATION:
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                                                                                                                                                                                              , Application US/08880179 6091004
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Similarity 93.3%;
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Friedrich, Leslie
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Pred. No. 63;
O; Mismatches
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Pred. No. 63;
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RESULT 12
US-08-972-927-2/c
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Best Local S
Matches 13
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REGISTRATION NUMBER: 38,241
REFERENCE/DOCKET NUMBER: CGC
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8587
TELEFAX: (919) 541-8689
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 9919 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 2, Application No. 6166290
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GENERAL INFORMATION:
APPLICANT: Rea, P
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, V.

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/880,179

FILING DATE:

CLASSIFICATION: 800
                                                           COUNTRY: US
ZIP: 19103-7086
ZIP: 19103-7086
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vo
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/972,927
FILING DATE: 18-NOV-1997
CLASSIFICATION: 800
CLASSIFICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ANTI-SENSE: 08-880-179-1
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TITLE OF INVENTION: PLANTS
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
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          PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 18-NOV-
PRIOR APPLICATION DATA:
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STREET:
CITY: Ph
STATE: P
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13; Conservative
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Lu, Yu-Ping
Li, Ze-Sheng
Li, Ze-Sheng
VENTION: GLUTATHIONE-S-CONJUGATE TRANSPORT IN
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18-NOV-1997
                       JMBER: US 60/031,040
18-NOV-1996
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Pred. No
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                                                                                                                                Version
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1.7e+02;
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                            TELEFAX: 617 876-5851
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 213 base pairs
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MOLECULE TYPE:
8-972-927-2
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APPLICANT: CELESTE, Anthony J.
TITLE OF INVENTION: BMP-11 COMPOSITIONS
NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 02140
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                      NAME: LAZAR, Steven R.
REGISTRATION NUMBER: 32,618
REFERENCE/DOCKET NUMBER: GI5205-A
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617 876-1170
TELEFAX: 617 876-5851
                                                                                                                                                                                                                                                              CLASSIFICATION:
ATTORNEY/AGENT IN
NAME: LAZAR, S
                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/247,907A
FILING DATE: May 20, 1994
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ADDRESSEE: (
                                                                      MOLECULE TYPE: |
ORIGINAL SOURCE:
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                                          STRAIN:
                                                         ORGANISM:
                                                                                                                TYPE: nuclei
                                                                                                    TOPOLOGY:
            NAME/KEY:
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3. 5639638
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GENETICS INSTITUTE, INC.
CambridgePark Drive
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V: 435
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Pred. No. 1.7e+02;
D; Mismatches 0;
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; NAME/KEY: misc_feature; LOCATION: 184..185; OTHER INFORMATION: /not; OTHER INFORMATION: of pus-08-452-772-3
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US-08-452-772-3/c
                                                                                                                                                                                                                                       CURRENT AFFLICATION NUMBER: US/08/452,
APPLICATION NUMBER: US/08/452,
FILING DATE: 30-MAY-1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/247
FILING DATE: 20-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: LAZAR, Steven R.
REGISTRATION NUMBER: 32,618
REFERENCE/DOCKET NUMBER: G152
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617 876-1170
TELEPHONE: 617 876-5851
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 213 base pairs
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Matches
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OTHER INFORMATION: /note OTHER INFORMATION: of page 8-247-907A-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: WOZNEY, John
APPLICANT: CELESTE, Anthony J.
TITLE OF INVENTION: BMP-11 COMPO
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: GENETICS INSTITUTE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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ORIGINAL SOURCE:
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ACTERISTICS:
LYPE: nucleic acid
STRANDEDNESS: do...
TOPOLOGY:
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LOCATION:
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CITY: Cambridge
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  . Similarity 13; Conserv
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US-09-414-234-3/c
; Sequence 3, Application
; Patent No. 6340668
; Patent INFORMATION:
; GENERAL INFORMATION:
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SEQUENCE DESCRIPT
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APPLICATION NUMBER: US/09/414,234
FILING DATE: 07-Oct-1999
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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ADDRESSEE: GENETICS INSTITUTE,
STREET: 87 CambridgePark Drive
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REGISTRATION NUMBER: 31,544
REFERENCE/DOCKET NUMBER: G152
ECOMMUNICATION INFORMATION:
TELEPHONE: 617 876-1170
TELEFAX: 617 876-5851
ION FOR SEQ ID NO: 3:
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iilarity 92.9%;
Conservative
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CELESTE, Anthony J.
THIES, R. Scott
INVENTION: BMP-11 COMPOSITIONS
F SEQUENCES: 11
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[N: Human A
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SECRIPTION: SEQ
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INFORMATION: /not
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2: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
3: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
4: /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
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6: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
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10: /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
11: /cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
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US-09-826-508-1
US-09-867-701-1577
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US-09-998-598-1600
US-09-738-626-252
US-10-037-182-17
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Sequence 1488, Ap
Sequence 1577, Ap
Sequence 1600, Ap
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	α	quence 14852,	equence 2	equence 2363	equence 46/,	ce 327,	quence 7	e 34920	Sequence 92/, App	ence 1978,	Sequence 8, Ap	e 167	quence	equence 16,	Ø	e 9041	equence 1	e 4348	Sequence 2	1	quence 1/5, A	Sequence 5109	equence 4484,	΄,	Sequence 2255	quence 95238,

ALIGNMENTS

RESULT 1
US-10-083-168-17/c
; Sequence 17, Application US/10083168
; Publication No. US20030023069A1
; GENERAL INFORMATION:

APPLICANT:
APPLICANT:
APPLICANT:

Liaw, Chen W. Chalmers, Derek T.

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RESULT 2
US-10-083-168-86/c
; Sequence 86, Applicat
; Publication No. US200
; GENERAL INFORMATION:
; APPLICANT: Liaw, Che
                                                                                                                                                                                                                                                                                                                      APPLICANT: Behan, Dominic P.
APPLICANT: Maciejewski-Lenior, Dominique
APPLICANT: Leonard, James N.
APPLICANT: Crtuno, Daniel
APPLICANT: Lin, I-Lin
TITLE OF INVENTION: Endogenous And No. US20(
TITLE OF INVENTION: Receptors
FILE REFERENCE: AREN-0320
CURRENT APPLICATION NUMBER: US/10/083,168
CURRENT FILING DATE: 2002-02-26
NUMBER OF SEQ ID NOS: 102
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Best Local S
Matches 14
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APPLICANT:
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TYPE: DNA
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Behan, Dominic
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                                                         Sequence 207, Applica Publication No. US200 GENERAL INFORMATION: APPLICANT: Behan, Doi APPLICANT: Chalmers
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TITLE OF INVENTION: Endogenous A
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CURRENT APPLICATION NUMBER: US/1
CURRENT FILING DATE: 2002-02-26
NUMBER OF SEQ ID NOS: 102
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TITLE OF INVENTION: No. US20030105292A1-Endogenous,
TITLE OF INVENTION: Protein-Coupled
TITLE OF INVENTION: Receptors
FILE REFERENCE: AREN-0040
CURRENT APPLICATION NUMBER: US/10/251,385
CURRENT FILING DATE: 2002-09-20
PRIOR APPLICATION NUMBER: US/09/170,496
PRIOR FILING DATE: 1998-10-13
NUMBER OF SEQ ID NOS: 294
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  REFERENCE:
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14; Conser
                                                        Behan, Domi
Chalmers,
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No. US20030105292A1
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AREN-0040
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Ners, Derek T.
                                                          Dominic P. ers, Derek T.
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RESULT 6
US-09-738-626-1488
; Sequence 1488, Application US/09738626
; Sequence No. US20020197605A1
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; ORGANISM: Homo
US-10-251-385-207
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US-09-826-508-1/c
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Best Local S
Matches 14
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SOFTWARE: F
SEQ ID NO 1
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APPLICANT: Nabil El
APPLICANT: Lisa Va
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SEQ ID NO 207
· LENGTH: 1446
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TITLE OF INVENTION: G Protein-Coupled Receptor Polypeptides
TITLE OF INVENTION: and Polynucleotides
FILE REFERENCE: GP-70744USB
CURRENT APPLICATION NUMBER: US/09/826,508
CURRENT FILING DATE: 2001-04-05
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CURRENT FILING DATE: 2002-09-20
PRIOR APPLICATION NUMBER: US/09/170,496
PRIOR FILING DATE: 1998-10-13
NUMBER OF SEQ ID NOS: 294
CURRENT APPLICATION NUMBER: US/09/738,626
CURRENT FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: JP 99/377484
PRIOR FILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: JP 00/159162
PRIOR FILING DATE: 2000-04-07
                                                                       APPLICANT: OZAKI, AKIO
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REFERENCE: 249-125
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TYPE: DA
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IKEDA,
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99;
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; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(390)
; OTHER INFORMATION: n = P
US-09-867-701-1577
                        GENERAL INFORMATION:

APPLICANT: Ruvkun, Gary

APPLICANT: Ogg, Scott

TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC TO

FILE REFERENCE: 00786/351004

CURRENT APPLICATION NUMBER: US/09/205,658

CURRENT FILING DATE: 1998-12-03

EARLIER APPLICATION NUMBER: 08/857,076

EARLIER APPLICATION NUMBER: 08/888,534

EARLIER APPLICATION NUMBER: US/09/205

EARLIER FILING DATE: 1997-07-07

EARLIER FILING DATE: 1997-07-07

EARLIER FILING DATE: 1998-05-15
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US-09-867-701-1577
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APPLICANT: Jones, Robert

APPLICANT: Harlocker, Susan L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CA

FILE REFERENCE: 210121.497

CURRENT APPLICATION NUMBER: US/09/867,701

CURRENT FILING DATE: 2001-05-29

NUMBER OF SEQ ID NOS: 10912
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LENGTH: 390
TYPE: DNA ORGANISM: Homo saf
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Matches 14
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Matches 1
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LENGTH: 375
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                                                                                                                                                                                                                                       Sequence 316, Application Patent No. US20010029617A1
NUMBER OF SOFTWARE:
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Similarity 93.3%;
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Pred. No. 2.2e+02;
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APPLICANT: Stolk, John A.

APPLICANT: Xu, Jiangchun

APPLICANT: Chenault, Ruth A.

APPLICANT: Meagher, Madelein Joy

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER

FILE REFERENCE: 210121.561

CURRENT APPLICATION NUMBER: US/09/998,598

CURRENT FILING DATE: 2001-11-16

CURRENT FILING DATE: 2001-11-16

TOWART OF SEQ ID NOS: 2606
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; ORGANISM: Caenorhabditis elegans
US-09-205-658-316
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US-09-998-598-1600/c
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US-09-998-598-1600
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US-09-738-626-
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APPLICANT: SENOH, AKIHIRO
APPLICANT: IKEDA, MASATO
APPLICANT: OZAKI, AKIO
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REFERENCE: 249-125
CURRENT APPLICATION NUMBER: US/09/738,626
CURRENT FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: JP 99/377484
PRIOR APPLICATION NUMBER: JP 00/159162
PRIOR FILING DATE: 2000-04-07
PRIOR APPLICATION NUMBER: JP 00/280988
PRIOR FILING DATE: 2000-08-03
NUMBER OF SEQ ID NOS: 7059
SOFTWARE: Patentin ver. 3.0
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SEQ ID NO 1600
LENGTH: 593
TYPE: DNA
ORGANISM: Hom
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Patent No. US200201
GENERAL INFORMATION
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OCHIAI, F
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Pred. No. 2.2e+02;
); Mismatches 1
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; LENGTH: 1350
; TYPE: DNA
; ORGANISM: Corynebacterium glutamicum
US-09-738-626-252
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US-10-044-090-438
                             ; FEATURE:
; NAME/KEY: CDS
; LOCATION: (1)..(4716)
US-10-037-182-19
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US-10-037-182-19
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                                                                                                                                                          APPLICANT: Thyboll, Jill
TITLE OF INVENTION: Recombinant Laminin 10
FILE REFERENCE: 99-274-F
CURRENT APPLICATION NUMBER: US/10/037,182
CURRENT FILING DATE: 2001-12-21
PRIOR APPLICATION NUMBER: 60/257,449
PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/279,282
PRIOR APPLICATION NUMBER: 60/279,282
PRIOR FILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 36
                                                                                                                                                                                                                                                                                                             Sequence 19, Applicat Publication No. US200 GENERAL INFORMATION: APPLICANT: Tryggvaso APPLICANT: Doi, Mas
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APPLICANT: Olga Bandman
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LENGTH: 7263
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Best Local
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LENGTH: 3239
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                                                                                                                                                  SOFTWARE:
                                                                                      ORGANISM: Mus musculus
                                                                                                        TYPE: DNA
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OTHER INFORMATION: Incyte ID No. US20020137081A1 241624.1
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Similarity 93.3%;
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Pred. No. 2.2e+02;
Mismatches 1;
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SOFTWARE: Patentin Ver. 2.

SEQ ID NO 17

LENGTH: 7554

TYPE: DNA
ORGANISM: Mus musculus
FEATURE:
NAME/KEY: CDS
LOCATION: (193)..(5007)
NAME/KEY: sig_peptide
COCATION: (193)..(291)
US-10-037-182-17
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                                                                                                                ; ORGANISM: HOMO US-09-764-868-1483
                                                                                                                                                                APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, P:
FILE REFERENCE: PTZ32
CURRENT APPLICATION NUMBER: US/09/76;
CURRENT FILING DATE: 2001-01-17
Prior application data removed - references
NUMBER OF SEQ ID NOS: 1510
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1483
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US-10-037-182-17
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Best Local S
Matches 14
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CURRENT FILING DATE: 2001-12-21
PRIOR APPLICATION NUMBER: 60/257,449
PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/279,282
PRIOR FILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 36
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TITLE OF INVENTION: Recombinant Laminin 10
FILE REFERENCE: 99-274-F
                                                                                                                                           TYPE: DNA
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Pred. No. 2.3e+02;
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RESULT 15
US-09-738-626-1/c
US-09-738-626-1/c

Sequence 1, Application US/09738626

Publication No. US20020197605A1

GENERAL INFORMATION:
APPLICANT: NAKAGAWA, SATOSHI
APPLICANT: HAYASHI, MIKIRO
APPLICANT: HAYASHI, KEIKO
APPLICANT: TATEISHI, NAOKO
APPLICANT: IKEDA, MASATO
APPLICANT: IKEDA, MASATO
APPLICANT: OZAKI, AKII

TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REFERENCE: 249-125
CURRENT APPLICATION NUMBER: US/09/738,626
CURRENT FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: JP 99/377484
PRIOR FILING DATE: 2000-04-07
PRIOR FILING DATE: 2000-08-03
NUMBER OF SEQ ID NOS: 7059
SOFTWARE: PatentIn ver. 3.0

LENGTH: 3309400
TYPE: DNA
ORGANISM: Corynebacterium glutamicum
US-09-738-626-1
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Search completed: June 26, Job time: 75.2 secs
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Best Local Similarity 93.3%;
Matches 14; Conservative
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BQ757051 EBem10_SQ
BM376561 EBem05_SQ
AJ434075 AJ434075
BQ461961 HD02L15r
BI777123 EBro03_SQ
BQ468011 HR01K11r
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ALIGNMENTS

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1601027 60324950	E427382 PSR6374	J034888 BJ034888	037202 ВJ037	1604112 6032440	J033194 ВJ033194	F530127 60204030	2432572 1M0218K0	G910143 6028055	J260486 ВJ260486	J302594 BJ30259	Q853560 QGB20N1	1669564 60329428	Q273893 rc72c10	773000 EST25410	I805258 S034E11	M027631 CsaC3001	F146084 WHE1837	29578 AL829578	E415681 MWL038.B	1950768 HVSME100	F623261 HVSMEa00	G418413 HVSMEk00	F254300 HVSMEf00	1954106 HVSMEm00	H446702 BOGVJ04	G415434 HVSMEk00	3415523 HVSMEk00	M099080 EBpi05_S	2471432 HV02G15r	2465677 HU04F22	H474182 BOGVM88	V915369 AV915369	2464764 HU01G04	Q471255 HV01N14	2465771 HU04K09	V941160 AV94116	1780309 EBem06_S	BG300356 HVSMEb001

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RESULT 1
BQ757051/c
LOCUS
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JOURNAL
COMMENT
                                                                                                                                                                                                                                                                                                                                                                                                                  ACCESSION
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KEYWORDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DEFINITION
                                        FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM
                       source
                                     Hordeum vulgare.

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooideae; Triticeae; Hordeum.

2E 1 (bases 1 to 237)

RS Hedley,P., Liu,H., Caldwell,D., McCallum,N., Mudie,S., Cardle,L., Ramsay,L., Machray,G., Marshall,D.F.M. and Waugh,R.

Development of Barley Transcriptome Resources

AL Unpublished (2001)

Contact: Waugh R, Marshall DF
Genome Dynamics/Computational Biology
Scottish Crop Research Institute
Invergowrie, Dundee, DD2 5DA, Scotland, UK
Tel: 00 44 1382 562731

Fax: 00 44 1382 562426

Taxil. Ast@scri.sari.ac.uk.
                                                                                                                                                                                                                                                                                                                                                                                                                                        BQ757051 . 237 bp mRNA linear EST 26-JUL-200; EBem10_SQ002_K07_R embryo, 2 Day germination, no treatment, cv Optic, EBem10 Hordeum vulgare cDNA clone EBem10_SQ002_K07 5', mRNA sequence. BQ757051 BQ757051.1 GI:21965523 EST.
                   est@scri.sari.ac.uk.
Location/Qualifiers
1. .237
/organism="Hordeum vulgare"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EST 26-JUL-2002
                                                                                                                                                                                                                                                                                                                                                             Pooideae
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RESULT 2
BM376561/c
·LOCUS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hedley, P., Liu, H., Caldwell, D., McCallum, N., Mudie, S., Ramsay, L., Machray, G., Marshall, D.F.M. and Waugh, R. Development of Barley Transcriptome Resources Unpublished (2001)
On Jan 10, 2002 this sequence warring
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EST.
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Hordeum vulgare
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooidea
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Scottish Crop Research Institute
Invergowrie, Dundee, DD2 5DA, Sco
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Tel: 00 44 1382 562731
Fax: 00 44 1382 562426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: est@scri.sari.ac.uk
All sequence has a Phred quality
Seq primer: M13 reverse.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
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/db_xref="taxon:4513"
/clone="EBem05_SQ002_J05"
/clone_lib="embryo, 14 DPA, rEBem05"
                 Ø
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0, 2002 this sequence
Waugh R, Marshall Dr
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/lab_host="DH10B"
/note="Vector: pSPORT1; Site_1: Sal I; Site_2: Not I;
/note="DH10B"
/note="DH
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/db_xref="taxon:4513"
/clone="EBem10_SQ002_K07"
/clone_lib="embryo, 2 Day germination, no
Optic, EBem10"
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-ras 0;
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VERSION KEYWORDS SOURCE

ORGANISM

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Matches

ORIGIN BASE COUNT

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence.
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AJ434075
 Molecular
Institute
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                                                                                                                                          sequence
BQ461961
BQ461961
EST.
                                                                                                                                                                                        BQ461963
HD02L153
       Unpublished (2002)
Contact: Stein Nils
Molecular Markon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P.O.Box Finland.
                                                                                         Hordeum vulgare
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooidea
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Similarity 100.0%;
15; Conservative 0
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooidea
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56 (Viikinkaari 6A),
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$00007 Hordeum vulgare
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/clone="S0000700014H03F1"
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/dev_stage="Shoot"
/note="2-,3-,4-days after germination"
117 c 116 g 100 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
Markers Group, Department Genbank of Plant Genetics and Crop Plant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (2002)
                                                                                                                                                        GI:21269732
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                                                                                 Hordeum.
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Mismatches 0;
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                                                                                                                                                                                         bp mRNA clone
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cDNA clone S0000700014H03F1,
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                                                                                                                                                                                         linear EST HD02L15 5-PRIME,
                                                                                                                                                                                                                                                                                                                                 Length 441;
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 Research (IPK)
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                                                          and Graner, A.
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mRNA
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BI777123/c
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                                                                                                 source
                                                                                                                                                Hedley, P., Liu, H., Caldwell, D., McCallum, N., Mudie, S., (Ramsay, L., Machray, G., Marshall, D.F.M. and Waugh, R. Development of Barley Transcriptome Resources
Unpublished (2001)
On Sep 26, 2001 this sequence version replaced gi:15780
Contact: Waugh R, Marshall DF
Genome Dynamics/Computational Biology
Scottish Crop Research Institute
Invergowrie, Dundee, DD2 5DA, Scotland, UK
Tel: 00 44 1382 562731
Fax: 00 44 1382 562426
Email: est@scri.sari.ac.uk
All sequence has a Phred quality score of 20 or over
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Insert Length: 'Plate: 2 row: 1
Seg primer: M131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: stein@ipk-gatersleben.de
Insert Length: 444 Std Error:
Plate: 2 row: L column: 15
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Tel: 039482-552
Fax: 039482-559
                                                                                                                                                                                                                                                                                                                                                                                                                                                         EBro03_SQ002_A18_R root, 3 week, waterlogged, cv Optic, EBro03 Hordeum vulgare cDNA clone EBro03_SQ002_A18 5', mRNA sequence.

BI777123
BI777123.2 GI:21948259
EST.
Hordeum vulgare.
Hordeum vulgare
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooidea; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooidea;
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primer: M13 reverse.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Hordeum vulgare"
/cultivar="Golden Promise"
/db_xref="taxon:4513"
/clone="HD02L15"
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/dev_stage="callus (5-10 mm in diameter)"
/lab_host="xL10-Gold"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone_lib="HD
/tissue_type="
/organism="Hordeum vulgare"
/cultivar="Optic"
/db_xref="taxon:4513"
/clone="EBro03_SQ002_A18"
/clone_lib="root, 3 week, w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             . 444
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. 7.5e+02;
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   Optic,
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BQ468011/c
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BQ468011
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HR01K11r
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Email:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EST sequencing and analysis in barley (2002) Unpublished (2002)
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15; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Molecular Markers Group, Department Genbank
Institute of Plant Genetics and Crop Plant Research (IPK)
Corrensstr. 3, 06466, Gatersleben, Germany
                                                                                                                                                                                                                                                                                                                                                                                                       Insert Length: 474
Plate: 1 row: K of Seq primer: M13rev
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Tel: 039482-5522
Fax: 039482-5595
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hordeum
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                   114
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                                                                                                                                                                                                                                                                                                                                                                                                                              ength: 474
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     a; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
phyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooidea
eae; Hordeum.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /tissue_type="root"
/dev_stage="3 week"
/lab_host="DH10B"
/note="Vector: pSPORT1; Site_1: Sal I; Site_2: Not I;
/note="Vector: pSPORT1; Site_1: Sal I; Site_2: Not I;
Non-normalised library, directionally cloned into pSPORT1.
Derived from roots of 3 week old waterlogged barley.
plants. Developed as part of the barley transcriptome resources of BBSRC/SEERAD funded cereal IGF (Investigating Gene Function) project."

Gene Function) project."
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                                /note="Vector: pBluescript SK+; Site_1: EcoRI (5'-end of cDNA); Site_2: XhoI (3'-end of cDNA); Due to a cloning artefact caused by the kit, in most cases the EcoRI site is NOT present, as well as the EcoRIadapter used for cloning. To excise the insert, restriction sites upstream EcoRI should be used (e.g. BamHI, SalI, PstI). NOTE: Also due to the cloning system used Blue/white selection for recombinats is not 100% reliable.Average insert size is 800 bp"
                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
1. .474
                                                                                                                                                                                                                                                                                             /organism="Hordeum vulgare"
/cultivar="Barke"
/db_xref="taxon:4513"
/clone="HR01K11"
                                                                                                                                                                                                   /tissue_type="roots"
/dev_stage="grown on filter paper at 25 Grad C
hours in petridish, 1-1.5 cm in length"
/lab_host="XL10-Gold"
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lumn: 11
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Query Match Best Local

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Score Pred.

15; No.

DB 14; 7.6e+02;

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Matches

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RESULT

DB 12; . 7.8e+02; ches 0;

Indels

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HVSMEb0012021f Hordeum vulgare s
HVcDNA0002 (Dehydration stress)
HVSMEb0012021f, mRNA sequence.
BG300356
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BG300356.1
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Clemson University Genomics Institute
Clemson University
100 Jordan Hall, Clemson, SC 29634, U
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864 656 4293
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Location/Qualifiers
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                                                                                                               conditions in the dark at room temperature on filter paper with water, nystatin and cefotaxime in covered crystallization dishes. Five-day old seedlings were incubated at 90% RH for 24 hr. Shoots were then harvested, total RNA was prepared, poly(A) RNA was purified, one primary unamplified cDNA library was made, 600000 pfu were in vivo excised to give pBluescript SK(-) cDNA phagemids. These steps were performed in the TJ Close laboratory at the University of California, Riverside (Choi, Close, Fenton). Phagemids were plated and picked at the Clemson University Genomics Institute (CUGI) (Begum, Palmer, Frisch, Atkins and Wing). Plasmid DNA preparations, DNA sequence nas been trimmed to remove vector sequence and contains a minimum of 100 bases of phred value 20 or above. For more details on library preparation and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rwing@clemson.edu
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http://www.genome.clemson.edu/projects/barley. To this clone see http://www.genome.clemson.edu/order see Close TJ, Wing R, Kleinhofs A, Wise R (2001) Genetically and physically anchored EST resources
                                                                      sequence analysis http://www.genome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone_lib="Hordeum vulgare seedling shoot EST library HVcDNA0002 (Dehydration stress)" /tissue_type="Seedling shoot" /lab_host="TJC121" /note="Vector: lambdaZAP; Site_1: EcoR1; Site_2: Xho1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seeds were
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/cultivar="Morex"
/db_xref="taxon:4513"
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Email: est@scri.sari.ac.uk
Email: est@scri.sari.ac.uk
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Dynamics/Computational Biology
h Crop Research Institute
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vulgare cDNA clone EBem06_SQ001_E02 5', mRNA sequence.
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/cultivar="Optic"
/db_xref="taxon:4513"
/clone="EBem06_SQ001_E02"
/clone_lib="embryo, 21 DPA, referred."
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/lab_host="DH10B"
/note="Vector: pSPORT1; Site_1: Sal I; Site_2: Not I;
/note="Vector: pSPORT1; Site_1: Sal I; Site_1: Not I;
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/note="Vector: pSPORT1; Site_1: Sal I; Site_1: Sal I;
/note="Vector: pSPORT1; Site_1: Sal I;
/note="Vector: pSPORT1; Site_1: Sal I;
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Sato, K., Saisho, D. ....
Barley EST sequencing project
Barley EST sequencing project
Unpublished (2002)
Contact: Tadasu Shin-i
Center For Genetic Resource Information
National Institute of Genetics
1111 Yata, Mishima, Shizuoka 411-8540, Ja
Tel: 81-559-81-6856
Fax: 81-559-81-6855
Fax: 81-559-81-6855
                                    Barley ESTs from germinating seeds
Unpublished (2002)
Contact: Stein Nils
Molecular Markers Group, Department Genbank
Institute of Plant Genetics and Crop Plant Res
Corrensstr. 3, 06466, Gatersleben, Germany
Tel: 039482-5522
Fax: 039482-5595
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Hordeum vulgare subsp. spontaneum
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. Similarity 100.0%;
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooidea;
Triticeae; Hordeum.
1. (bases 1 to 565)
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Email: stein@ipk-gatersleben.de Insert Length: 565 Std Error: Plate: 4 row: K column: 9
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/strain="H602"
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/clone_lib="K. Sato unpublished cDN
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/tissue_type="top three leaves"
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aceae; Pooideae
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RESULT 11
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DEFINITION
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AUTHORS
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15; Cons
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BQ471255.
EST.
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Zhang,H.,
                                                                                                                                                                                                                     Email:
Insert
Plate:
Seq pri
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1 (bases 1 to 575)
                                                                                                                                                                                                                                                                                       Corrensstr. 3, 06
Tel: 039482-5522
Fax: 039482-5595
                                                                                                                                                                                                                                                                                                                                                                           Contact
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/lab_host="XL10-Gold"
/note="Vector: pBluescript SK+; Site_1: EcoRI (5'-end of cDNA); Site_2: XhoI (3'-end of cDNA); germinating seeds, 16-48h. Due to a cloning artefact caused by the kit, in most cases the EcoRI site is NOT present, as well as the EcoRIadapter used for cloning. To excise the insert, restriction sites upstream EcoRI should be used (e.g. BamHI, SalI, PstI). NOTE: Also due to the cloning system used Blue/white selection for recombinats is not 100% reliable. Average insert size is 1.2 kb"
                                                                                                                                                                                                                                                      stein@ipk-gatersleben.de
Length: 575 Std Error:
                                                                                                  /organism="Hordeum vulgare"
/cultivar="Barke"
/db_xref="taxon:4513"
/clone="HV01N14"
/clone_lib="HV"
                                                                                                                                                                                                                                                                                                                       r Markers Group, Department Genbank
e of Plant Genetics and Crop Plant Research (IPK)
tr. 3, 06466, Gatersleben, Germany
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HV Hordeum vulgare cDNA clone HV01N14 5-PRIME, mRNA
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/cultivar="Barke"
/db_xref="taxon:4513"
/clone="HU04K09"
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/note="Vector: pBluescript SK+; Site_1: EcoRI (5'-end of cDNA); Site_2: XhoI (3'-end of cDNA); Roots were grown for two days on filter paper at room temperature. Due to a
                                                                                                                                                                                        Location/Qualifiers
1. .575
                                                 /tissue_type="germinating seeds"
/dev_stage="germinating seeds (48-96 h)"
/lab_host="XL10-Gold"
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Pred. No. 8.1e+02;
o. mismatches 0;
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RESULT 12
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                                     Match
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Similarity 100.0%;
15; Conservative
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Tel: 039482-5522
Fax: 039482-5595
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BQ464764 597 bp mRNA linear EST HU01G04T HU Hordeum vulgare cDNA clone HU01G04 5-PRIME,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Stein Nils
Molecular Markers Group, Department Genbank
Institute of Plant Genetics and Crop Plant Research
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                                                                                         49
                                                                                                                                                                                                                                                                                                                                                                                                                                                       stein@ipk-gatersleben.de
Length: 597 Std Error:
1 row: G column: 4
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138 c 159 g 138 t
                                                                                    /note="Vector: pBluescript SK+; Site_1: EcoRI (5'-end of cDNA); Site_2: XhoI (3'-end of cDNA); germinating seeds, 16-48h. Due to a cloning artefact caused by the kit, in most cases the EcoRI site is NOT present, as well as the EcoRIadapter used for cloning. To excise the insert, restriction sites upstream EcoRI should be used (e.g. BamHI, SalI, PstI). NOTE: Also due to the cloning system used Blue/white selection for recombinats is not 100% reliable. Average insert size is 1.2 kb"
                                                                                                                                                                                                                                                                                                                /organism="Hordeum vulgare"
/cultivar="Barke"
/db_xref="taxon:4513"
/clone="HU01G04"
/clone_lib="HU"
                                                                                                                                                                                                                                                            /tissue_type="germinating seeds"
/dev_stage="germinating seeds, 16-48 h"
/lab_host="XL10-Gold"
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   0;
Score 15; DB Pred. No. 8.2; Mismatches
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Pred. No. 8.1e+02;
Mismatches 0;
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DB 14;
8.2e+02;
thes 0;
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                                  Length 597;
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BASE COUNT
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AV915369/c
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Barley EST so
Unpublished
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15; Cor
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Center F
                                            Whole genome shotgun sequencing of Unpublished (2001) Other GSSs: BOGVM88TR Contact: Chris Town
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Fax: 81-
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Fax: 301-838-0208
Email: cdtown@tigr.org
DNA is from a doubled haploid
Seq primer: TF
Class: sheared ends.
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Eukaryota; Viridiplantae; Liliopsida; Poales; Poaceae; Pooidea
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooidea
; Triticeae; Hordeum
1 (bases 1 to 607)
Nichalek, W., Weschke, W., Stein, N. and Grane
                                                                                                                                                                                                                                                                                                                                                                                  Barley ESTs from germinating seeds
Unpublished (2002)
Contact: Stein Nils
Molecular Markers Group, Department Genbank
Institute of Plant Genetics and Crop Plant 1
Corrensstr. 3, 06466, Gatersleben, Germany
Tel: 039482-5522
Fax: 039482-5595
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/tissue_type="germinating seeds, 16-48 h"
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/lab_host="XL10-Gold"
/note="Vector: pBluescript SK+; Site_1: EcoRI (5'-end of cDNA); Site_2: XhoI (3'-end of cDNA); germinating seeds, 16-48h. Due to a cloning artefact caused by the kit, in most cases the EcoRI site is NOT present, as well as the EcoRIadapter used for cloning. To excise the insert, restriction sites upstream EcoRI should be used (e.g. BamHI, SalI,PstI). NOTE: Also due to the cloning system used Blue/white selection for recombinats is not 100%
                                                                                                                                                                                                                                                                                                                                stein@ipk-gatersleben.de
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/strain="TO1000DH3"
/db_xref="taxon:3712"
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Copyright (c) 1993 - 2003 Compugen Ltd.
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Direct Submission

Submitted (10-FEB-1994) Ian Tomlinson, MRC Centre for Prendineering, Hills Road, Cambridge, CB2 2QH, U.K

3 (bases 1 to 165)

Cook, G.P., Tomlinson, I.M., Walter, G., Riethman, H., Carte Buluwela, L., Winter, G. and Rabbitts, T.H.

A map of the human immunoglobulin VH locus completed by the telomeric region of chromosome 14q

Nat. Genet. 7 (2), 162-168 (1994)

95004581

7920635
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mammalia; Metazoa; Chordata; C:
Mammalia; Eutheria; Primates; C:
1 (bases 1 to 165)
Cook,G.P., Tomlinson,I.M., Walte-
Buluwela,L., Winter and D-1
Unpublished
2 (base-
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(DA-10).
Z29985
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Similarity 100.0%;
15; Conservative (
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4-1VH3,
         Homo sapiens
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llarity 100.0%;
Conservative (
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                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
/isolate="der 8 chromosome f
/db_xref="taxon:9606"
/dev_stage="adult"
/germline
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4 c 3 g
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/codon_start=1
/product="immunoglobulin
(DA-10)"
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36 c 44 g
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         217 bp
rearranged
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Pred. No. 1.4e+03;
                                                                                                                           Score 15; DB 9; Pred. No. le+03; Mismatches
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Rabbitts,T.H.
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Catarrhini;
         immunoglobulin heavy
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                    PRI 10-MAR-1998
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TITLE
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2 (bases I to

Delabie, J. and Tierer

Delabie, J. and Tierer

Delabie, J. and Tierer

Delabie, J. and Tierer
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Homo sapiens
Eukaryota; Met
Mammalia; Euth
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variable regic
AF017465
AF017465,1 G1
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marginal
Unpublish
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Mammalia; E
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AJ227738.
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Similarity 100.0%;
15; Conservative (
                                                                                                                                                           Direct Submission
Submitted (08-AUG-1997) Pathology, University of Leuven,
Minderbroedersstraat 12, Leuven B-3000, Belgium
Location/Qualifiers
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                                                                                           /isolate="case 5"
/db_xref="taxon:9606"
<1. .>223
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Minderbroedersstraat 12, B-3000, BELGIUM
                                                                                                                                                                                                                                                                                          iens
                                                                      /gene="IgH"
<1. .>223
                                                                                                                                                                                                                                                                                                                                                       223 bp mRNA linear PRI 18-OCT-19
iens isolate case 5 rearranged immunoglobulin heavy chain
region (IgH) mRNA, partial cds.
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/product="immunoglobulin heavy chain variable region"
/protein_id="CAA12716.1"
/db_xref="GI:2951654"
/translation="FTFSDYYMSWIRQAPGKGLEWLSDMSSSGTSIYYADSVKGRFTI
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             /codon_start=2
/product="rearranged immunoglobulin heavy chain variable
region"
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55 c 59 g 45 t
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/protein_id="AAB81258.1"
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                                                       'gene="IgH"
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and Tierens, A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15; DB 9; Pred. No. 1e+03; Mismatches 0
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Matches
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Best Local S
Matches 15
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ORGANISM
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          variable region.
AJ306240
AJ306240.1 GI:1264174
2c-18m3 gene; immunoql
                                                                                                                                                                                                                                                                                                                                                     Evidence for c marginal zone Unpublished
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Homo sapiens
Eukaryota; Met
Mammalia; Euth
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Similarity 100.0%;
15; Conservative
                                                                                                                   20
                                                                                                                                                                                                                                                                                                                                                                                   Direct Submission
Submitted (03-MAR-1998) Delabie J., Pathology, University
Leuven, Minderbroedersstraat 12, B-3000, BELGIUM
2 (bases 1 to 223)
Delabie, J. and Tierens, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens 7-9VH3.
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Similarity 100.0%;
15; Conservative
human.
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sapiens
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/translation="SGFTFSDYYMSWIRQAPGKGLEWVSYISSSSSYTNYADSVKGRF
TISRDNAKNSLYLQMNSLRAEDTAVYYCAR"
54 c 59 g 49 t
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Eutheria;
1 to 223)
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/product="immunoglobulin heavy chain variable region"
/protein_id="CAA12712.1"
/db_xref="GI:2951646"
/translation="SGFTFSDYYMSWIRQAPGKGLEWVSYISSSSSYTNYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCAR"
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<1. .>223
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54 c 59 g 4
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immunoglobulin;
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Primates;
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rearranged
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No. le+03;
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; Hominidae;
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Saarlandes,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thesis (2000) Department of Medizinische Klinik,
Universitaetskliniken des Saarlandes, Homburg, Germany
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Mammalia;
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15; Cons
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Eukaryota;
Mammalia; E
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                                                                                                                                                                                                                                                                                                                                                             Homo sapiens clone 5-4.3.10A immunoglobulin heavy chain variable region gene, partial cds.
                                                                      Submitted (18-AUG-2000) Internal Medicine and Harold C. Simmons Arthritis Research Center, University of Texas Southwestern Medical Center, 5323 Harry Hines Blvd., Dallas, TX 75235, USA

Location/Qualifiers
1. .255
                                                                                                                                                                                      Lee, J. and Lipsky, P.E.
The V lambda J lambda repertoire in human fetal spleen:
for positive selection and extensive receptor editing
Unpublished
                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                              Lee, J. and Lipsky, P.E. Direct Submission
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/product="immunoglobulin heavy chain variable region"
/product= 68 g 54 t
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<1. .>246
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/db_xref="taxon:9606"
/tissue_type="Synovial membrane"
/rearranged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (06-JAN-2001) Weisgerber K.,
s, Onkologisches Labor, 66424
Location/Qualifiers
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/db_xref="taxon:9606"
/chromosome="14"
/clone="5-4.3.10A"
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                                                            organism="Homo
                                             'isolate="FS5"
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Pred. No. 9.9e+02;
Mismatches 0;
                                                          sapiens"
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Mammalia; Eutheria; Primates; C
1 (bases 1 to 258)
Foerste, N., Gellrich, S., Golemb
Sterry, W. and Jahn, S.
Analysis of VH genes rearranged
dermal infiltrates of patients
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jahn, S.
Direct Submission
Submitted (21-MAR-1997) S. Jahn, Univers
Department of Dermatology, Schumannstrasse
Location/Qualifiers
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Y12032
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V-region.
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                  /gene="EMZ112VH3"
/codon_start=1
/product="immunoglobulin heavy chain"
/product="immunoglobulin heavy chain"
/protein_id="CAA72763.1"
/db_xref="GI:1914733"
/translation="SGFSFSDYYMSWIRQAPGKGLDWVSYISDNSNFRYSADSVKGRFTISRDNSKNSVFLHMSSLRADDTAVYYCARGRTAPRGPEDYW"
1 66 c 67 g 59 t
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/codon_start=1
/product="immunoglobulin heavy chain virg/product="immunoglobulin heavy chain virg/product="
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<1. .>255
                                                                                                                                                                                                                                                                                                      /dev_stage=
/rearranged
                                                                                                                                                                                                                                                                                                                           /cell_type="B lymp.
/tissue_type="skin
/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
/isolate="patient EM"
/db_xref="taxon:9606"
/chromosome="14"
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/tissue_type="spleen
/dev_stage="fetus; l
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Pred. No.
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Catarrhini;
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9.8e+02;
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; Hominidae; Homo.
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20/21, 10117 Berlin,
                                                                                                                                                                                                                                                                                 fungoides
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRI 27-MAR-1997 (EMZ112VH3).
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Direct Submission
Submitted (21-SEP-:
Frankfurt, Theodor
Homo sapiens partial IGVH3-8 gene for variable region, donor MF, cell4. AJ415895 GI:16117163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              centers a
Blood 97
21093792
11157489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleosi
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 261)
Brauninger, A., Yang, W., Wacker, H.H., Rajewsky, K., Kuppers, R. Hansmann, M.L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            عا
                                                                                                                                         9
                                                                                                                                                                                                 Similarity
15; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B-cell development in progressively transformed germinal centers: similarities and differences compared with classical germinal centers and lymphocyte-predominant Hodgkin disease Blood 97 (3), 714-719 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HSA406644 261 bp DNA linear PRI 22-MAR Homo sapiens partial VH3-11 gene for rearranged immunoglobulin heavy chain variable region, isolate case4-PTGC1-171.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        immunoglobulin heavy chain; variable region; VH3-11 gene
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larity 100.0%;
Conservative (
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rity 100.0%;
nservative 0
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                                                                                                                                                                                                                                                                                                                                /codon_start=1
/product="immunoglobulin heavy chain variable region"
/protein_id="CAC10708.1"
/db_xref="GI:10636713"
/translation="SGFTFSDYYMSWIRQAPGKGLEWVSYISSSGSTIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCARDVRPLWLHYFDYW"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (21-SEP-2000) Brauninger A., Theodor Stern Kai 7, 60590, Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                        /gene="VH3-11"
<1. .>261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/isolate="case4-PTGC1-171"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                        1. .>261
/gene="VH3-11"
/product="immunoglobulin heavy
/product="71 g 62 t
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                                                                                                                                                                                                                                                                                                                                                                                                                       'gene="VH3-11"
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Pred. No. 9.8e+02;
Mismatches 0;
                                               immunoglobulin heavy chain
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                                                                                                                                                                                                                                                                                        chain variable
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Eur. J. Immunol.
2 (bases 1 to 273)
Goossens, T., Brauninger, A., k
Receptor revision plays no ma
repertoire of human memory B
hypermutation
Eur. J. Immunol.
3 (bases 1 to 273)
                                                                                                                                               Homo sapiens
Eukaryota; Metazoa; (
Mammalia; Eutheria;
1 (bases 1 to 278)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Brauninger, A.

Direct Submission
Submitted (21-SEP-2001) Brauninger A.,
Frankfurt, Theodor Stern Kai 7, 60590,
                                                                                                1 (bases 1 to 278)
Kanzler,H., Kueppers,R., Hansmann,M.L.
Hodgkin and Reed/Sternberg cells in Holoutgrowth of a dominant tumor clone degerminal center B cells
                                                                                                                                                                                                                                                     gene (
277321
                                                                                                                                                                                                                                                                                                                                                                                                        Similarity 100.0%; 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Brauninger, A., Goossens, T., Rajewsky, K. and Ku
Regulation of immunoglobulin light chain gene
early B cell development in the human
Eur. J. Immunol.
2 (bases 1 to 273)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Mammalia; Eutheria; 1 (bases 1 to 273)
Direct Submission
Submitted (18-JUL-1996) Kanzler
University of Cologne, Weyertal
3 (bases 1 to 278)
Kanzler, H., Kuppers, R., Hansmann
                                                                                                                                                                                                                                          277321.1
                                                                                                                                                                                                                                                                 H.sapiens gene for immunoglobulin heavy gene (V3-11).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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                                                                                     Unpublished
                                                                                                                                                                                                                 region
                                                                                                                                                                                                                            immunoglobulin heavy
                                                               Kanzler, H.
                                                                                                                                                                                                      Homo
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<1. .>273
/gene="IGVH3-8"
/product="immunoglobulin he
/product="78 g 62 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /isolate="donor MF"
/db_xref="taxon:9606"
/clone="cell4"
/cell_type="B cell"
/rearranged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /gene="IGVH3-8"
<1. .>273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="naive lambda expressing
1, .273
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Location/Qualifiers
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                                                                             278)
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Primates;
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Primates;
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                                                                                                                                                                                                                             chain;
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MF"
 Hansmann, M.L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  major role in shaping the B cells after the onset
                                                                                                                                                                                                                             immunoglobulin superfamily;
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                                                                                                                                                                 Craniata; Vertebrata;
Catarrhini; Hominidae;
                         H.,
121,
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Hodgkin's disease represent
derived from (crippled)
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; Hominidae;
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 Rajewsky, K.
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the receptor
                          Genetics
1, FRG
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J. Exp. 1
97033523
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AF077461
AF077461
                                                                                                                                                                                                                                                           Lipsky,P.
Direct Su
Submitted
Center, 5
                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota;
Mammalia;
1 (bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hodgkin
outgrowt
                                                                                                                                                                                                                                                                                                                 Brezinschek, H.P., Dorner, T., Farner, N.L., Brezinschek, R.I. and Lipsky, P.E.
The Influence of CD40-CD154 Interactions on the Expressed Human Variable Heavy Chain Repertoire: Analysis of VH Genes Expressed by Individual B Cells of a Patient with X-Linked Hyper-IgM Syndrome Unpublished

2 (bases 1 to 279)
Brezinschek, H.P., Dorner, T., Farner, N.L., Brezinschek, R.I. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9
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                                                                                                                                                                                                                                                                                         Submission
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rity 100.0%;
nservative 0
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                                                                                                                                                        /db_xref="Homo sapiens"
/db_xref="taxon:9606"
/chromosome="14"
/clone="BF3N2g3E09"
                                                                                                                                                                                                                                                          d (23-OCT-1998) Internal 5323 Harry Hines Blvd., I
/note="VH(D)JH"
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STIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCARDIGVWG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
1. .278
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'cell_type="Hodgkin-RS"
'tissue_type="Hodgkin's lymphoma"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /isolate="patient 8"
/db_xref="taxon:9606"
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                                                                                                                                      rearranged
                                                                                                                      'note="from patient with
                                                                                                                                                      'cell_type="individual B
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Pred. No. 9.7e+02;
Mismatches 0;
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Dallas, TX
                                                                                                                        X-linked hyper-IgM syndrome"
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VERSION
KEYWORDS
                           RESULT 14
AY003824
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AY013309
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AY013309
AY013309.1 GI:17220530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (08-NOV-2000) Internal Medicine and Harold C. Simmons Arthritis Research Center, University of Texas Southwestern Medical Center, 5323 Harry Hines Blvd., Dallas, TX 75235, USA Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lee, J. and Lipsky, P.E. Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished 2 (bases 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mammalia; Eutheria; Primates; Catarrh 1 (bases 1 to 279)
Lee, J. and Lipsky, P.E.
VH gene replacement in human fetal B immunoglobulin heavy chain receptor e
   HOMO
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Eukaryota; M
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Similarity 100.0%;
L5; Conservative
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                                                                                                                                        Conservative
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/db_xref="taxon:9606"
/chromosome="14"
/clone="FM1-3.3.12A"
/cell_type="IgM+ B lymphocyte"
/tissue_type="bone marrow"
/dev_stage="fetal; 18 weeks gestation"
/rearranged
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                                                                                                                                                                                                      /note="CDR2 replacement"
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ADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCARDHYGSGSFDYW"
67 c 77 g 65 t
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283 bp
sy-3M/9-H1
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Pred. No.
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Pred. No. 9.7e+02;
Mismatches 0;
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Comparative characteristics of mu chain and alpha chain transcripts expressed by individual tonsil plasma cells
Mol. Immunol. 38 (1), 19-34 (2001)
11483207
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AY003824
AY003824.
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Hillson, J.
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Mammalia;
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15; Cor
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Yavuz,S.
                                                                                                                                   and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Direct Submission Submitted (02-JUN-2000) Autoimmunity Branch, National Institute of Arthritis and Musculoskletal and Skin Diseases, 9000 Rockville Pike Bldg. 9 1N124, Bethesda, MD 20892, USA
                                                 Hillson, J.L., Karr, N.S., Oppliger, I.R., Mannik, M. and Sasso, E.H. The structural basis of germline-encoded VH3 immunoglobulin binding to staphylococcal protein A J. Exp. Med. 178 (1), 331-336 (1993)
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                                                                                                           Immunology (sites)
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                                                                                                                       ison, J.L., Oppliger, I.R., Sasso, E.H., Milner, E. and Wener, M.H emerging B cell repertoire: Influence of developmental stage interindividual variation
Immunology 149, 3741-3752 (1992)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sapiens.
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                                                                                                                                                                                                                                     n; immunoglobulin heavy chain V-region.
piens fetus liver cDNA to mRNA.
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nservatíve
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/cell_type="plasma cell
<1. .>283
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1. .283
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/db_xref="GI:15777398"
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NAKNSLYLQMNSLRAEDTAVYYCATDYSGYNRVLLELYWGQGTLVTVSSG"
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Location/Qualifiers
1. .291
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Nanki,T., Yavuz,A.S. and Lipsky,P.E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note="VH3 family, 3-11/DP-35"
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Eutheria; Primates; Catarrhini;
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Pred. No. 9.7e+02;
Mismatches 0;
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linear

PRI 06-JAN-1995

Homo.

Indels

0

Gaps

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BASE COUNT
ORIGIN
Search completed: June Job time : 427.6 secs
                                                                                                               Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 15; Conservative
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/cell_line="L2E5"
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/gene="IGHV@"

/note="CDR-2; G00-

175. .270

/gene="IGHV@"

/note="framework 3

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/gene="IGHV@"

/note="CDR-3; G00-

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/note="CDR-1;
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/note="This CDS feature is included to show the
/note="This CDS feature is included to show the
/rotein of the corresponding V_region. Presently
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/gene="IGHV@"
/product="immunoglobulin heavy chain
/note="G00-128-528"
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Adams CW, Carter PJ, Fendly BM, Gurney AL; WPI; 1999-204666/17.		25-AUG-1997; 97US-0918148.	21-AUG-1998; 98WO-US17364.	04-MAR-1999.	WO9910494-A2.	Homo sapiens.		neuromuscular; muscular dystrophy; complementarity determining region; variable heavy chain; variable light chain; VH; VL; ss.	; myelotoxic chemotherapy; leukaemia;	oplasia; disseminated intravascular coag	yocyte; platelet; immunological; hematopoieti	throm		Ah2 variable heavy (VH) chain CDR1 encoding DNA.	17-JUN-1999 (first entry)		222303.	RESULT 1 AAX32393 ID AAX32393 standard; DNA; 15 BP.

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RESULT 2
AAZ46860
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             increase platelet production. They can be used for treating immunological or hematopoietic disorders, especially thrombocytopenia. Thrombocytopenia -associated bone marrow hypoplasia (e.g. aplastic anem following chemotherapy or bone marrow transplant) may be effectively treated with the antibody compounds as well as disorders such as disseminated intravascular coagulation (DIC), immune thrombocytopenia (HIV-induced and non HIV-induced), chronic idiopathic thrombocytopenia, congenital thrombocytopenia, thrombotic thrombocytopenia and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       used for improving neuromuscular function in a patient, e.g. in muscular dystrophy. The products can also be used for detection and diagnosis. The antibodies have a longer half-life than the natural ligand for the TPO-R. Sequences AAX32387-X32413 represent DNA fragments encoding the CDR1, CDR2, and CDR3 regions of variable heavy (VH) chains and variable light (VL) chains of antibodies Ab1 to Ab6.
                                                                                                                                      07-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  myelodysplasia. They can also treatment of solid tumours or autologous or allogeneic bone idiopathic aplastic anemia, co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (TPO). They can stimulate proliferation, differentiation or growth o megakaryocytes. They may also be able to stimulate megakaryocytes to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         thrombopoietin
be used in the
(TPO). They can
WPI; 2000-116276/10
                            Cramer DV,
                                                                                                                                                                                                   W09958157-A1
                                                                                                                                                                                                                                                                                                                           Human immunoglobulin IgHV3-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to
                                                          (CRAM/) CRAMER D V. (KEAR/) KEARNS-JONKER
                                                                                                         08-MAY-1998;
                                                                                                                                                                       18-NOV-1999
                                                                                                                                                                                                                                                                   anti
                                                                                                                                                                                                                                                                                                                                                            10-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ46860 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     idiopathic aplastic anemia, congenital thrombocytop
thrombocytopenia. The antibodies which bind to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New thrombopoietin
                                                                                                                                                                                                                                                                                              Anti-xenogeneic
                                                                                                                                                                                                                                                               -amyloid
                                                                                                                                                                                                                                  sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   . Similarity 15; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ention relates to an agonist antibody (Ab) which binds to oletin receptor (TPO-R). The antibodies which bind the in the same way and for the same indications as thrombothey can stimulate proliferation, differentiation or growthey can stimulate proliferation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
illarity 100.0%;
Conservative 0
                                                                                                                                                                                                                                                                                 disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               B₽;
                            Kearns-jonker
                                                                                                                                                                                                                                                            antibody; Alzheimer's
                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                          98US-0084725
                                                                                                                                       99WO-US09971.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 A; 4 C;
                                                                                                                                                                                                                                                                                              antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 86pp;
                                                                                                                                                                                                                                                                               systemic
                                                                                                                                                                                                                                                                                                                                                                                                                        294 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    bone marrow transplant, myelodysplasia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English
                                                                                                                                                                                                                                                                                              antibody;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  or agonist antibodies - hematological disorders
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                                                                                                                                                                                                                                                                                                                          Vh gene sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                be used in e.g. myelotoxic chemothera leukaemia, myeloablative chemotherapy
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Т;
                                                                                                                                                                                                                                                              xenograft rejection; diab
erythematosus; rheumatoid
disease; or anti-insulin;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     thrombocytopenia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  useful
                                                                                                                                                                                                                                                                               rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MuSK receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      as thrombopoietin
                                                                                                                                                                                                                                                                                              diabetes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chemotherapy
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ne TPO-R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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The invention provides new methods for preventing expression of anti-
C xenogeneic antigen antibodies. The method comprises (i) selecting genes,
mRNA, or proteins associated with the expression or regulation of
C antibodies to xenogeneic tissue, organ or antigens; (ii) obtaining ONs
C which are associated with expression or regulation of the
C which are associated with expression or regulation of expression of the
C antibodies or aromatic amino acid oligomers comprising an amino acid
C sequence selected from antibody fragments targeted to minor microgrooves
C of the polynucleotides encoding them having at least one aromatic amino
C acid, and (iii) administering the ONs obtained to a subject to bind to
C its target and prevent, reduce or inhibit expression of anti-xenogeneic
C antigen antibody. The methods can be used for preventing, reducing or
C inhibiting xenograft rejection. The xenograft may be e.g. liver, heart
C and kidney organs, or islets of Langerhans tissues. The methods can also
C be used for treating autoimmune diseases or conditions e.g. systemic
C lupus erythematosus, rheumatoid arthritis, anti- amyloid antibodies
(Alzheimer's disease) or anti-insulin auto-antibodies (diabetes). The
products can also be used for detection of anti-xenogeneic antigen
antibodies and nucleic acids encoding them. The present sequence
C represents a human immunoglobulin IgHV3-11 Vh gene sequence binding to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Use of ar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Use of antisense oligonucleotides for preventing xenograft rejectitreating autoimmune conditions, e.g. systemic lupus erythematosus, rheumatoid arthritis, Alzheimer's disease or diabetes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P-PSDB;
                                                target
                                                   nucleic
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                                                   acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  34pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           rejection
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ф Query Match Best Local S Matches 15 Sequence 91 GACTACTACATGAGC 105 15; Con: \vdash GACTACTACATGAGC 15 294 BP; 68 A; 72 C; servative 100.0%; 0; 90 G; Score 15; DB 21; Pred. No. 65; Mismatches 0. 64 T; 0 other; 0 Length 294; Indels 0; Gaps 0;

RESULT 3 ABA94229 CB2iE7 ABA94229 standa 13-MAR-2002 ABA94229; heavy ch (first entry) ain variable region nucleotide sequence. rd; DNA; 329 BP.

Neuromodulatory ebvHigM MsI19D1 antiparkinsonian; central nervous system; CNS; sHIgM22; LYM 22; AKJR4;
; ebv HIgM CB2bG8; CB2iE12; CB2iE7; MSI19E5; virucide;
; neuroprotective; nootropic; vulnerary; ds.

Homo sapiens.

Location/Qualifiers 1..329 /*tag= /transl_exc /note= "Xaa עם _except= "(pos: 1..2; aa: xaa)" 'xaa = unknown"

WO200185797-A1

15-NOV-2001.

30-MAY-2000; 20)00WO-US14902

10-MAY-2000; $\frac{2}{2}$)00US-0568351.

(MAYO-) MAYO FOUND MEDICAL EDUCATION RES

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RESULT 4
AASO3478
ID AASC
XX
AC AASC
XX
DT 29-P
XX
DE DNA
XX
KW Anti
KW hear
XX
PN WO2C
XX
PN WO2C
XX
PD 19-P
XX
PR 11-C
XX
PA (CAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention provides a neuromodulatory agent (I) capable of promoting neurite outgrowth, regeneration, remyelination and neuroprotection in central nervous system (CNS). (I) is capable of inducing remyelination, promoting cellular proliferation of glial cells, and promoting Ca2+ csignaling with oligodendrocytes. An humanised antibody to (I) can be selected from antibody sHIGM22 (LYM 22), ebvHigM MsI19D10, ebv HIGM CE2bG8, AKJR4, CB2iE12, CB2iE7 or MsI19E5. (I) is useful for stimulating cells in consumer of cNS axons, or treating demyelinating proliferation of glial cells in consumer of consumer of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            neuroprotection post-infectious
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 promoting neurite
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rodriguez
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             where nerves are damaged as by trauma. The present the CB2iE7 heavy chain variable region nucleotide s
                                                                                                                                                                               Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                Antibody; adip heart disease;
                                                                                                                                                                                                                                                                                                                                          AAS03478;
                                                                                                                                                                                                                                                                                                                                                                            AAS03478
                                                                                                                                             WO200127279-A1
                                                                                                                                                                                                                                                                                                        29-AUG-2001
                                     12-OCT-1999;
                                                                        11-OCT-2000;
                                                                                                           19-APR-2001.
   (CAMB-)
                                                                                                                                                                                                                                                                   encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2002-066596/09
)B; ABB07182.
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                                                                                                                                                                              sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
   CAMBRIDGE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GACTACTACATGAGC
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                                                                                                                                                                                                                                                                                                                                                                             standard;
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                                                                                                                                                                                                                                 adipocyte;
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                                                                                                                                                                                                                                                                    anti-adipocyte monoclonal antibody
                                                                                                                                                                                                                                                                                                      (first
                                                                         2000WO-GB03900.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Latory agent (a human IgM monoclonal antibody), te outgrowth, regeneration, remyelination and in central nervous system, useful to treat encephalomyelitis -
                                                                                                                                                                                                                cyte; heavy chain; light chacomplementarity determining
                                       99US-0158812.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              81
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     ANTIBODY TECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                             cDNA;
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                                                                                                                                                                                                                                                                                                        entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   80
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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66;
                                                                                                                                                                                                                                   chain;
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                                                                                                                                                                                                                                                                    heavy chain,
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RESULT 5
AAL43586
ID AAL4
XX
AC AAL4
XX
DT 05-9
XX
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Best Local S
Matches 15
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ligand
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and light chain
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                                                                                                                                                                       Key
CDS
                                                                                                                                                                                                                                                           26-10
                                                                                                                                                                                                           Unidentified.
                                                                                                                                                                                                                                                                                  Dig3 antibody heavy chain coding sequence.
                                                                                                                                                                                                                                                                                                                                                             AAL43586
                                                                                                                                                                                                                                                                                                            05-SEP-2002
                                                                        02-MAY-2002
                                                                                               WO200234886
                       27-OCT-2000;
                                              26-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          light chain coding sequences of the invention. The antibody heavy id in the treatment of obesity and obesity related diseases. The cibodies can be used to deliver drugs or pro-drugs directly to the fat self. Antibodies binding specifically to adipocytes can be used to ivate the immune system to destroy the cells by complement mediated is. The antibodies may be labeled with a detectable label such as iiolabel, fluorescent or chemical group and used in methods of ignosis in human subjects e.g. to determine the presence of adipocyte sence or level of adipocytes in a cell or tissue sample. The ibodies can be used as an alternative means of treatment for obese iionts other than undergoing surgery to remove excess fat. Antibodies of different types of fat deposits can also be produced e.g. intra-lominal fat associated with heart disease.
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B; AAU02578
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binding pr
d structure
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                       2000US-0699023.
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 XAS SYSTEM.
                                                                                                                                                                                                                                                                                                            irst entry)
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                                                01WO-US46795.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /29.
                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  binding members of antibody molecules which bind to is used in the treatment of obesity and obesity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              141; 182pp; English.
                                                                                                                       /product= "Dig3 heavy chain"
/note= "No start or stop codon is
                                                                                                                                               /partial
                                                                                                                                                            /*tag=
                                                                                                                                                                                                                                                                                                                                                               DNA;
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RESULT
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AS03470
AAS03470
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Best Loo
Matches
                                                                           WPI; 200
P-PSDB;
                         Panel of specific whole adipocytes related diseases
   Disclosure;
                                                                                                                 Edwards
                                                                                                                                                                 12-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention comprises a method of obtaining a bacterium containing a nucleic acid encoding a binding protein capable of binding a target ligand, or a nucleic acid encoding a catalytic protein that catalyses a reaction involving a target substrate. The method of the invention allows the isolation of a cell expressing a nucleic acid encoding a binding protein capable of binding a target ligand, and consequently the isolation of the nucleic acid. The binding protein produced by the method of the invention is useful for the de novo isolation of antibodies from large repertoire libraries and for monitoring production of a particular byproduct of a biological reaction. The present DNA sequence encodes the dig3 scFv antibody heavy chain which is used in an example of the invention.
                                                                                                                                                                                         11-OCT-2000;
                                                                                                                                                                                                                  19-APR-2001
                                                                                                                                                                                                                                           WO200127279-A1
                                                                                                                                                                                                                                                                    Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Match
                                                                                                                                                                                                                                                                                        rt disease; complementarity de
                                                                                                                                                                                                                                                                                                                               encoding
                                                                                        2001-
                                                                                                                                                                                                                                                                  sapiens
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AAO15187.
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 Page
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                                                                                                               Main
                                                                                                                                                                                                                                                                                                                             anti-adipocyte
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        вP;
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                                     c binding is used i
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  36;
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                                                                                                                                        ANTIBODY
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182pp;
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                                     members of ant in the treatmen
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                                                                                                                                       TECHNOLOGY
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Englis
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Pred.
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No.
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tment of c
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                                                                                                                                                                                                                                                                                                     chain;
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66;
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                                                                                                                                                                                                                                                                                       obesity; ion; CDR;
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                                                                                                                                                                                                                                                                                                                          heavy chain,
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                                   and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           351;
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                                                                                                                                                                                                                                                                                                                             FAT
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В

AC XXX DIT XXX OS XXX O

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This sequence en (CDR) from an an specific binding antibody antigen of less than 1.0

encodes a VH domain complementarity determining region entibody specific for estradiol. The invention relates to g members (sbp) comprising a polypeptide that comprises an binding domain (AABD) which has a dissociation constant 0x10^-8M for estradiol, and a dissociation constant of at

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RESULT
AAZ3165
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antigen on the presence or leve antibodies can patients other for different t
                                                                                                                                                                                                                                                                                                                                                                    antibody antigen
dihydrotestoster
menstrual cycle;
diagnosis; VH do
                                                                                               estradiol
therapy an
                                                                                                                                                  WPI; 19:
P-PSDB;
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itself. Antibo
activate the i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            lysis. The an radiolabel, f diagnosis in
                                                                         Disclosure;
                                                                                                                                                                                        Johnson
                                                                                                                                                                                                                                                                                            02-NOV-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Coding
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                                                                                              or levels
                                                                                                                                                                                                                                                                  1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence
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abel, flu
                                                                                                                                                                                                               CAMBRIDGE ANTIBODY TECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antibodies binding specifically to adipocytes can be used as a therapeutic Antibodies binding specifically to adipocytes can be used to be the immune system to destroy the cells by complement mediated. The antibodies may be labeled with a detectable label such as abel, fluorescent or chemical group and used in methods of the surface of an adipocyte to detect or determine the sis in human subjects e.g. to determine the presence of adipocyte or level of adipocytes in a cell or tissue sample. The lies can be used as an alternative means of treatment for obeses other than undergoing surgery to remove excess fat. Antibodies iferent types of fat deposits can also be produced e.g. intra-lal fat associated with heart disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity .5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                    complementarity determining region; CDR; estriol-3-sulphate; ntigen binding domain; steroid hormone; estriol; testosterone; tosterone; progesterone; estriol 3-beta-di-glucuronide; cycle; hormone replacement therapy; oestrogen secreting tumour; VH domain; ds.
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                                                                                                                                                                                     Pope AR,
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                                                                                           nding partners for estradiol, used for monitoring s during the menstrual cycle, in hormone replacement diagnosing oestrogen secreting tumours
                                                                                                                                                              .3/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   535 represent anti-adipocyte monoclonal antibody heavy n coding sequences of the invention. The antibodies can be eatment of obesity and obesity related diseases. The be used to deliver drugs or pro-drugs directly to the fat
                                                                      umn 21-22; 26pp;
                                                                                                                                                                                                                                        96US-0028897.
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                                                                                                                                                                                                                                                                                                                                                                                                                                           for VH domain CDR of anti-estradiol antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA; 366 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               determining the presence or absence of estradiol in a sample. They can used for monitoring estradiol levels, e.g. during the menstrual cycle, hormone replacement therapy and for diagnosing oestrogen secreting tumours. The sbps can provide for discrimination between estradiol and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 200
P-PSDB;
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                                                                                                                                                          and light chain coding sequences of the invention. The antibody heavy used in the treatment of obesity and obesity related diseases. The antibodies can be used to deliver drugs or pro-drugs directly to the finance of an obese patient or the antibody can be used as a therapeutic itself. Antibodies binding specifically to adipocytes can be used to activate the immune system to destroy the cells by complement mediated lysis. The antibodies may be labeled with a detectable label such as radiolabel, fluorescent or chemical aroun and mand in the label such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antibody; adip heart disease;
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radiolabel, fluorescent or chemical group and used in methods of diagnosis in human subjects e.g. to determine the presence of adipantigen on the surface of an adipocyte to detect or determine the presence or level of adipocytes in a cell or tissue sample. The antibodies can be used as an alternative means of treatment for ol patients other than undergoing surgery to remove excess fat. Antil for different types of fat deposits can also be produced e.g. intrabolation and the deposition of the produced e.g. intrabolation and the deposition of the deposition also be produced e.g. intrabolation and the deposition of the deposition and the deposition of the deposition also be produced e.g. intrabolation and the deposition of th
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                                                                                                                                                                                                                                                                                                                                                                                                 English.
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to 5 familie
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                Lable heavy chains. The genes were isolated and cloned from a series cosmid constructs: Y202; Y103; Y21; Y6;Y24; 3-31; M84; M118 and M131, Y2R amplification using primers AAQ78917-38. The genes are subdivided that fragments were isolated from high molecular weight DNA fragments were isolated from high molecular weight DNA from placenta. The DNA was partially digested with TaqI restriction me. The fragments were separated by gel elecrophoresis and 35-45 kb constitutions were collected. The fragments were ligated with ClaI-digested nid vector pJB81. The ligation products were in vitro packed and cotted into E.coli 490A. The fragments were then subcloned by colony cidisation. The Vh genes and the DNA fragments encoding them are full in producing human immunoglobulin in mammalian hosts.
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human immunoglobulin in mammalian hosts
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252..348
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terminator or splice site sequence"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA; 715 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               90 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 15; DB Pred. No. 66; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               119 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunoglobulin; variable; heavy chain;
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                                                                                                                                                                                                                                                                                                                                                                                does not conform
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 369;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  heavy
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RESULT 10
AAC00031
ID AAC00031 s
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Best Local S
Matches 15
                                          Query Match
Best Local s
Matches 13
                                                                                                                 The present sequence is one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. An ORF has been identified within the sequence. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full lengt cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and secretion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo
                                                                                                                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-SEP-2000.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-OCT-2000
                                                                                                                                                                                                                                                                         Claim 1;
                                                                                                                                                                                                                                                                                           obtaining cDNAs and genomic diagnostic, forensic, gene 1
                                                                                                                                                                                                                                                                                                                 New nucleic
                                                                                                                                                                                                                                                                                                                                                                      Dumas Milne Edwards
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human;
                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                          (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                       PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        therapy;
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Similarity 100.0%;
15; Conservative
                                        Similarity 86.
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                                                                                                                                                                                                                                                                       SEQ ID 29; 71pp + CD-ROM;
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GACTACTACATGAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard;
                                                                                     379 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EST;
                                                                                                                                                                                                                                                                                                                 acid that is a
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3
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                                                                                     A; 88 C;
                                                     94.7%;
86.7%;
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Pred.
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                                                                                     102 G;
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                                          d. No. 1.9 Mismatches
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                                                               14.2;
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1.9e+02;
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                                                                                     21 other;
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l to 5'ESTs
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RESULT

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RESULT 12
AAQ78960
ID AAQ78
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Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                   mass of an obese patient or the antibody can be used as a therapeutic itself. Antibodies binding specifically to adipocytes can be used to activate the immune system to destroy the cells by complement mediated lysis. The antibodies may be labeled with a detectable label such as radiolabel, fluorescent or chemical group and used in methods of diagnosis in human subjects e.g. to determine the presence of adipocyte antigen on the surface of an adipocyte to detect or determine the presence or level of adipocytes in a cell or tissue sample. The antibodies can be used as an alternative means of treatment for obese patients other than undergoing surgery to remove excess fat. Antibodies for different types of fat deposits can also be produced e.g. intra-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAS03474 standard; cDNA; 363 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antibody; adip
heart disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAS03474;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and light chain coding used in the treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Panel of specific binding members of antibody molecules which bind to whole adipocytes is used in the treatment of obesity and obesity related diseases -
                                                                  AAQ78960;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      antibodies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-2820
P-PSDB; AAU025
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                                        03-AUG-1995
                                                                                            AAQ78960
                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                              abdominal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   encoding anti-adipocyte monoclonal antibody heavy chain,
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14; Cor
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                                                                                                                                                                                                                93.3%;
arity 100.0%;
onservative 0
                                                                                            lard;
             obulin Vh gene #22.
                                                                                                                                                                                                                                                                                              associated with heart disease
                                                                                                                                                                                                                                                                                                                                                                                                                                         535 represent anti-adipocyte monoclonal antibody heavy n coding sequences of the invention. The antibodies can be eatment of obesity and obesity related diseases. The be used to deliver drugs or pro-drugs directly to the fat se patient or the antibody can be used as a therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ge 139; 182pp; English.
                                         first entry)
                                                                                                                                                                                      CTACATGAG 14
                                                                                                                                                                                                                                                                    P; 84 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ain SH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DGE ANTIBODY TECHNOLOGY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cyte; heavy chain; light chain; obesity; FAT; complementarity determining region; CDR; ss.
                                                                                            DNA;
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                                                                                            ВP.
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Pred. No.
                                                                                                                                                                                                                                                                     103 G;
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). 2.4e+02;
0;
                                                                                                                                                                                                                                                                    T; 0 other;
                                                                                                                                                                                                                                          Length 363;
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ABK63583
ID ABK6
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AC ABK6
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DT 18~5
XX
DE Rat
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Best Local S
Matches 14
                                                                                                                                                                                                                                                                              A series of genes (AAQ78939-79002) encoding human immunoglobulin variable heavy chains. The genes were isolated and cloned from a series of cosmid constructs: Y202; Y103; Y21; Y6;Y24; 3-31; M84; M118 and M131, by PCR amplification using primers AAQ78917-38. The genes are subdivided into 5 families of Vh genes. The fragments cover a region of 800 kb. The DNA fragments were isolated from high molecular weight DNA from human placenta. The DNA was partially digested with TaqI restriction enzyme. The fragments were separated by gel elecrophoresis and 35-45 kb fractions were collected. The fragments were ligated with ClaI-digested cosmid vector pJB81. The ligation products were in vitro packed and infected into E.coli 490A. The fragments were then subcloned by colony hybridisation. The Vh genes and the DNA fragments encoding them are useful in producing human immunoglobulin in mammalian hosts.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 19
P-PSDB;
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cosmid;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-MAY-1993;
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  Rat sequence
                                                                                                                                                                                                                                                          Sequence
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                              18-JUN-2002
                                                        ABK63583
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                                                                                                                                                                                                     similarity 100
4; Conservative
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                                                                                  standard;
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enta; vector;
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   differentially
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545..54
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428..43
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131..233
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85..544
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                                                                                 CDNA;
                             entry)
                                                                                                                                                                                                                                                        A; 128
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expressed
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                                                                                                                                                                                                    Score 14; DB Pred. No. 2.5; Mismatches
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                                                                                                                                                                                                                                                        184 G; 144
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   'n.
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2.5e+02;
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hosts
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   hepatotoxin
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   #1490.
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Query Match Best Local S Matches 14

Similar 14; Con:

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100.0%;

Score 14; Pred. No

DB 24; L. 2.7e+02; 0;

Length 1422;

93.3%;

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Mismatches No.

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GACTACTACATGAG 14

Sequence

1422

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305 C;

246 G;

446 T;

0 other;

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expression in a tissue or cell sample exposed to the compound of two or more genes listed in the specification, where differential expression of the genes is indicative of at least one toxic effect or progression. The method can also be used to identify an agent which modulates the toxic response and predict cellular pathways that a compound modulates in a cell. The methods utilise a set of at least two probes (on a solid support in kit form), where each of the probes comprises a sequence that specifically hybridises to a gene listed in the specification, a computer system comprising a database containing information identifying the expression level in a tissue or cell sample exposed to a hepatotoxin of a set of genes comprising at least two genes listed in the specification, and for identifying the expression level in a tissue or cell of at least one gene listed in the specification of the information used to present information that the specification and for identifying toxicity markers in changes in gene expression and for identifying toxicity markers in this sues or cell exposed to a known toxin. The genes may be used as toxicity markers for the prediction or identification of the physiological state of tissue or cell sample that has been exposed to a compound or agent. Hepatotoxicity sequence is an expressed sequence tag (EST) or cDNA derived from a gene which is differentially expressed in response to a hepatotoxic agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-JUL-2000;
02-NOV-2000;
11-MAY-2001;
15-MAY-2001;
22-MAY-2001;
06-JUN-2001;
13-JUN-2001;
19-JUN-2001;
09-JUL-2001;
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effects by determining the
cells exposed to the toxin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seq ID
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2000US-244880P.
2001US-290029P.
2001US-290645P.
2001US-292336P.
2001US-295798P.
2001US-297457P.
2001US-298884P.
2001US-303459P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               of a compound,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       in gene expression in tissues or cells exposed to ring these to gene expression in unexposed tissues luded are methods of predicting at least one toxic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       elates to methods for predicting toxic effects of progression of these toxic effects by determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           toxin;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             239pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of compounds or the progression of these the changes in gene expression in tissues or in and comparing these to gene expression i
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    comprising detecting
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Matches 14;
                                                                                                                                                                                         The invention relates to a polypeptide having GT1209, GT1354 or GT0946 activity obtained from Arabidopsis. The invention may also be applied to the development of herbicide tolerant plants, plant tissues, plant seeds and plant cells. The polypeptide is useful as herbicide targets in screening assays to identify potential herbicides and inhibitors of GT1209, GT1354 or GT0946 activity. A compound having herbicidal activity is useful for suppressing the growth of a plant. The newly discovered GT1209, GT1354 or GT0946 genes are essential for seedling growth and development. The present sequence is Arabidopsis thaliana GT1209 gene
                                                                                                                                                                                                                                                                                                                                                             WPI; 200
P-PSDB;
         AAD31644;
                             AAD31644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Key
                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                                   New polypeptide having GT1209, GT1354 or GT0946 activity, obtain Arabidopsis, useful as herbicide targets in screening assays to identify the inhibitors or potential herbicides
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DB; AAE19876.
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                                                                                                                                                                                                                                                                                                       WPI; 20
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                                                                               Sequence 3236
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arity 100.0%;
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                                  0;
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2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
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US-08-471-044-18
US-08-471-046A-18
US-08-471-046A-18
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US-08-471-033-26
US-08-471-033-26
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US-08-471-044-26
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US-08-545-809A-11
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GENERAL INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/958,201
FILING DATE:
PRIOR APPLICATION DATA:
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FILING DATE: 21-OCT-1996
INFORMATION FOR SEQ ID NO: 5
INFORMATION FOR SEQ ID NO:
                           TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-5070
TELEFAX: 617-542-8906
                                                                                                                       APPLICATION NUMBER: PCT/JP93/00603
FILING DATE: 10-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                  OPERATING SYSTEM: Wind
SOFTWARE: FastSEQ for
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Honjo, Tasuku
APPLICANT: Matsuda, Fumihiko
APPLICANT: Matsuda, Fumihiko
TITLE OF INVENTION: HUMAN IMMUNOGLOBULIN VH GENE
TITLE OF INVENTION: SEGMENTS AND DNA FRAGMENTS CONTAINING THE SAME
NUMBER OF SEQUENCES: 145
                                                                                                                                                                    APPLICATION NUMBER: US/0. FILING DATE: 27-MAR-1996 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
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STREET: 22
CITY: Bost
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                                                                         NAME: Freeman, John W. REGISTRATION NUMBER: 29 REFERENCE/DOCKET NUMBER:
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GENERAL INFORMAT
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TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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FILING DATE:
PRIOR APPLICATION N
APPLICATION N
FILING DATE:
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LENGTH: 715 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Honjo, Tasuku
APPLICANT: Matsuda, Fumihiko
TITLE OF INVENTION: HUMAN IMMUNOGLOBULIN VH GENE
TITLE OF INVENTION: SEGMENTS AND DNA FRAGMENTS CONTAINING THE SAME
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REFERENCE/DOCKET NUMBER: 0650
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-5070
TELEFAX: 617-542-8906
TELEX: 200154
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STRANDEDNESS:
TOPOLOGY: li
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OPERATING SYSTEM:
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ZIP: 02110-2804
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human lymphoblast
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FastSEQ for Windows Version
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Fish & Richardson, P.C.
Franklin Street
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RESULT 5
US-09-221-017B-404
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В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 650-494-0792
TELEX: 706141
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 2073 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                    Query Match
Best Local
                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: PP2911
FILING DATE: 09-APR-1998
PRIOR APPLICATION NUMBER: PCT/AU98
FILING DATE: 10-DEC-1000
ATTORNEY/ACCORN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 31-DEC-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PP1546
FILING DATE: 30-JAN-1998
PRIOR APPLICATION
                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
HYPOTHETICAL:
ANTI-SENSE: UN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE MEDIUM TYPE: D
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APPLICATION NUMBER: US
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CORRESPONDENCE ADDRESS:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 32,430
REFERENCE/DOCKET NUMBER: 27
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Monroy, Gladys H
                                                                                                                                                                                                                                                                                                                                                          ORIGINAL
                                                                                                                                                                                                                                                                                                    FEATURE:
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CITY: 1
STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 2
CLASSIFICATION:
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                                                                                                                                                                                                                                              NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 650-494-0792
                                                                                                                                                                                                                                                                                                                            ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
 1828
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                                                       GACTACTACATGAG 14
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23-DEC-1998
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N: P. GINGIVALIS NUCLEOTIDES AND USES
ES: 1120
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                                                                                                                                      Score 14;
Pred. No
                                                                                                              Mismatches
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                                                                                                                                           No.
                                                                                                                                      DB
38;
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US-08-767-128-37
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US-08-767-128-
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Patent No.
GENERAL IN
  Best Log
                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 35,093
REFERENCE/DOCKET NUMBER: 8648.49USF1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 612/371-5278
TELEFAX: 612/332-9081
INFORMATION:
               Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 04-DEC-1996
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/09258
FILING DATE: 05-JUN-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/541,373
FILING DATE: 10-OCT-1995
PRIOR APPLICATION NUMBER: 08/462,798
APPLICATION NUMBER: 08/462,798
                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM Compain OPERATING SYSTEM: DO SOFTWARE: FastSEQ VOICEMENT APPLICATION DAY APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                            TOPOLOGY: li
MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
ATTORNEY/AGENT
NAME: Carter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NERAL INFORMATION:
APPLICANT: WYLIE, DWANE E.
APPLICANT: WYLIE, DWANE E.
APPLICANT: MYLIE, DWANE E.
APPLICANT: LOPEZ, OSVALDO
APPLICANT: MURRAY, PETER JOSEPH
APPLICANT: GOEBEL, PETER
TITLE OF INVENTION: LEAD BINDING POLYPEPTIDES AND
TITLE OF INVENTION: NUCLEOTIDES CODING THEREFORE
NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESSEE: MG
STREET: 3100
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MEDIUM TYPE:
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PRIOR APPLICATION DATA:
                                                                                                                                                     FRAGMENT
                                                                                                                                       ORIGINAL SOURCE:
                                                                                                                                                                    ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
CITY: 1
STATE:
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                                                                                                                                                                                                                                              LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                           NAME/KEY: Coding LOCATION: 1...372 OTHER INFORMATION:
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14; Con
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                                                                                                                                                                                                                                            : 372 base pairs nucleic acid
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                                                                                                                                                       TYPE:
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FastSEQ Version 1.5
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: Genomic DNA
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)0 No. 6111079west Center,
                                                                                                                                                                                                                                                                                                                                                                                               NUMBER: 08/462,798
05-JUN-1995
INFORMATION:
89.3%;
rrity 93.3%;
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                                                                                                                                                                                                                               single
                                                                                                          Sequence
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 Score 13.4; D
Pred. No. 69;
0; Mismatches
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                                 DB 3;
      1;
                                   Length 372;
       Indels
       0;
       Gaps
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GATTACTACATGAGC

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LOCATION: 1..2004; OTHER INFORMATION: /note: OTHER INFORMATION: sequer US-08-471-033-18
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US-08-471-033-18
                                     Query Match
Best Local
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                         Matches
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                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: CG
TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8582
TELEFAX: 919-541-8689
INFORMATION FOR SEQ ID NO: 18:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 23-MAR-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: CIBA-GEIGY Corporation
                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMA
NAME: Pace, Gary M.
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA: APPLICATION NUMBER: FILING DATE: 09-SEP-
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                                                                                                                                                                       MOLECULE TYPE:
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                                                                                                                                    FEATURE:
                                                                                                                                                            HYPOTHETICAL:
                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                 ANTI-SENSE:
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STATE:
                                                                                                                                                                                                                                                                                                                                               FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: CLEA CERTIFICATION OF THE PROPERTY TO SKYLING Drive
                                                                                                                                                                                TYPE: STRANDEDNESS: SI
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                         CENGTH:
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                          14;
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                       Similarity 93.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hawthorne
                                                                                                                                                                                                            nucleic acid
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  GACTACTACATGAGC
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                                                                                                                                                                                                                          2004
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Koziel,
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Nye, Gordon J
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                                                                                                                                                             Ö
                                                                                                                                                                                                                                                                                                                                    INFORMATION:
                                                                                                                                                                          DNA
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23-MAR-1994
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25-MAR-1993
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Michael
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                                   89.3%;
93.3%;
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                                                                                   /note= "Maize optimized DNA sequence for VIP1A(a) 80 kd
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ER: CGC
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                        Score 13.4; D
Pred. No. 82;
0; Mismatches
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                      Gaps
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RESULT 8
US-08-471-044-18
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                                    US-08-471-044-18
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            Query Match
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                                                                                                                                                                                                        TELEFAX: 919~54
INFORMATION FOR SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30B
                                                                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8582
TELEFAX: 919-541-8689
                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 05-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 800 PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
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APPLICATION NUMBER:
FILING DATE: 25-MAR-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER
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                                                                                              FEATURE:
                                                                                                          ANTI-SENSE:
                                                                                                                     HYPOTHETICAL:
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STATE:
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                                             NAME/KEY: misc_feature LOCATION: 1..2004 OTHER INFORMATION: /not OTHER INFORMATION: sequ
                                                                                                                                                                                                                                                                                                                                      FILING
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/471,044 FILING DATE: 06-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                      STRANDEDNESS
                                                                                                                                                                                                                                                      REGISTRATION NUMBER REFERENCE/DOCKET
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                                                                                  NAME/KEY:
                                                                                                                                              TOPOLOGY:
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Representation
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Duck, Nicholas
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                                                                                                           NO
                                                                                                                                            linear
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e, Gordon J
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25-MAR-1993
                                                                                                                                DNA (genomic)
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23-MAR-1994
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NUMBER: CGC
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Michael
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No.
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Michael G
Martha A
 89.3%;
93.3%;
                                              sequence
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                                              /note= "Maize optimized DNA sequence for VIP1A(a) 80 kd
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Score 13.4;
Pred. No. 82;
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            DB 2;
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OTHER INFORMATION: 05-08-463-483A-18
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US-08-463-483A-18
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                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 08/
FILING DATE: 23-MAR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 25-MAR-1993
                                                                                                                                                                                      TELEPHONE: 919-541-8615
TELEFAX: 919-541-8689
INFORMATION FOR SEQ ID NO:
                                                                                                          TOPOLOGY: 1i
MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Vo
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/463,483A
                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Spruill, W. Murray
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: CG
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                 ANTI-SENSE:
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Amner: 7 Skyline Drive
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Koziel,
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Desai, Nali
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Michael
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  89.3%;
93.3%;
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   13.4;
No. 82;
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             Length 2004;
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US-08-471-046A-18
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GENERAL
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                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/463,483

FILING DATE: 05-JUN-1995

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/314,594

FILING DATE: 09-SEP-1994

PRIOR APPLICATION NUMBER: US 08/218,018

APPLICATION NUMBER: US 08/218,018
                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8587
TELEFAX: 919-541-8689
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS
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APPLICATION NUMPER
FILING TOTAL
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                                                                               MOLECULE TYPE:
                                              FEATURE:
                                                                    HYPOTHETICAL:
                                                         ANTI-SENSE:
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CLASSIFICATION:
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OTHER
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                                                                                                      STRANDEDNESS
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                       LOCATION:
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o. 5866326
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'KEY: misc_feature
PION: 1..2004
RINFORMATION: /note
RINFORMATION: seque
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Nye, Gordon J
                                                                                                                                                                                                                                                                                                                                                                                                                         PatentIn Release #1.0, CATION DATA:
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ziel, Michael G
llins, Martha A
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                                                                                                                                                                                                          , J. Timothy NUMBER: 38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESS:
No. 5866326artis Corporation
Cornwallis Road
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25-MAR-1993
INFORMATION:
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uch, Juan J
                                                                              DNA (genomic)
                                                                                                     single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Protein Genes
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/note= "Maize optimized DNA
sequence for VIP1A(a) 80 kd protein from AB78"
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RESULT 11
US-08-470-566B-18
~~mence 18, Ap
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                                                                                                                                      TELEFAX: 919-54
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APPLICANT:
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC Computer:
                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Meigs, J. Timothy
REGISTRATION NUMBER: 38,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 25-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Estruch, Juan J
APPLICANT: Estruch, Juan J
APPLICANT: Estruch, Juan J
TITLE OF INVENTION: NO. 5872212el Pesticic
NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS;
CORRESPONDENCE ADDRESS;
CORRESPONDENCE No. 5872212artis Corporation
                           ANTI-
                                                                                                                     FORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8587
TELEFAX: 919-541-8689
                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 08/463,483 FILING DATE: 05-JUN-1995 PRIOR APPLICATION DATA:
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CITY: I
STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                  TOPOLOGY:
                                                                              TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 08/218,018 FILING DATE: 23-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 0 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/314,594 FILING DATE: 09-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
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                           SENSE:
 KEY:
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872212
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Koziel, Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kostichka, N.
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                           NO
misc_feature
                                                                  linear
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06-JUN-1995
                                                                             single
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                                                    (genomic)
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93.3%;
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Kristy
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No. 82;
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; OTHER INFORMATION:
; OTHER INFORMATION:
US-08-470-566B-18
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US-08-469-334-18
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Best Local
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                                                                                         TELEFAX: 91
INFORMATION FOR
                                                                                                                                                                                                                                                                 PRIOR APPLICATION: 800
APPLICATION DATA:
APPLICATION NUMBER: 08
FILING DATE:
APPLICATION
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PRIOR APPLICATI
APPLICATION N
                                                                                                                          REFERENCE/DOCTELECOMMUNICAT
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                      MOLECULE TYPE:
                                                                                                                                                             ATTORNEY/AGENT NAME: Sprui
                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 2004 base pair
                                                                                                                                                                                                                               FILING DATE: 09-SEP-PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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                                            STRANDEDNESS
                                                                                                              TELEPHONE:
                                                                                                                                                                                    FILING DATE:
                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP:
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NY
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                                              nucleic acid
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2: CIBA-GEIGY Corporation
7 Skyline Drive
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Carr, Brian
Desai, Nalini
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Koziel, Michael G
Mullins, Martha A
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NO
                                                                                      N NUMBER: 32,943
OCKET NUMBER: CGC
TION INFORMATION:
919-541-8615
19-541-8689
SEQ ID NO: 18:
                                                                                                                                                                                          TION DATA:
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           N
O
                                                                                                                                                                       JUMBER: US 08/037,057
25-MAR-1993
INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                           E FORM:
Floppy disk
                                                                                                                                                                                                                                                    MBER: US 08/314,594
09-SEP-1994
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                                          single
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No.
                     (genomic)
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sequence for VIP1A(a) 80 kd
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Pred. No. 82;
                                                                                                                                       1695/CIP3
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RESULT 13
US-09-300-529-18
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Best I
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CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6066783artis C
STREET: 3054 Cornwallis Road
CITY: Research Triangle Park
STATE: NC
COUNTRY: USA
                      TELEPHONE: 919-541-8587
TELEFAX: 919-541-8689
INFORMATION FOR SEO TO SEQUENCE
                                                                                                                                                                                                                                                                                                   FILING DATE: TBA
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 06-JUN-1995
PRIOR APPLICATION NUMBER: US 08/
FILING DATE: 05-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 09-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 23-MAR-1994
PRIOR APPLICATION DATA:
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NAME/KEY:
LOCATION:
                                                  INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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APPLICATION NUMBER: US
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                                                                                                                                                                                                                            APPLICATION NUMBER: US 0 FILING DATE: 25-MAR-1993 ATTORNEY/AGENT INFORMATION:
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14; Conservative
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: 3054 Cornwallis Road
Research Triangle Park
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CHARACTERISTICS: 2004 base pair
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ENTION: Genes
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                                                                                                                                                                                                                                                                                                                                                                                                        JMBER: US 08/314,594
09-SEP-1994
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hka, N.
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93.3<del>8</del>;
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Kristy
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ed. No. 82;
Mismatches
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VIP1A(a) 80 kd
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LOCATION: 1.2004
COTHER INFORMATION:
COTHER INFORMATION:
US-09-300-529-18
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US-08-471-033-35
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Best Local S
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Patent No
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MOLECULE TYPE:
HYPOTHETICAL:
          INFORMATION FOR SEQUENCE CHAR!
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                                                                                           FILING DATE:
ATTORNEY/AGENT
NAME: Pace,
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APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                       COMPUTER READAB MEDIUM TYPE:
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                          REFERENCE/DOCKET NUMBER: P-40,403
REFERENCE/DOCKET NUMBER: CGC
TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8582
TELEFAX: 919-541-8689
                                                                                                                                                                        CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/314,594
FILING DATE: 09-SEP-1994
PRIOR APPLICATION DATA:
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CITY: I
STATE:
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ADDRESSEE: CIBA-GEIGY Corporation
STREET: 7 Skyline Drive
                                                                                                                                                                                                                                                                      SOFTWARE:
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5770696
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INVENTION: No. 5770696el Pesticidal Proteins and Strains
F SEQUENCES: 50
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CHARACTERISTICS: 2576 base pair
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Desai, Nalini M
Kostichka, N. K
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: Floppy disk
IBM PC compatible
YSTEM: PC-DOS/MS-DOS
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ziel, Michael
llins, Martha
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                                                                                Gary M.
NUMBER:
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single
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                                                                                                                 MBER: US 08/037,057
25-MAR-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nicholas B
                         ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gregory W
Michael G
Martha A
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93.3%;
pairs
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sequence for VIP1A(a) 80 kd
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Pred. No. 82;
0; Mismatches
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                                                                                                                                                                                                                                                                          Version
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; NAME/KEY: CDS; LOCATION: 9..2564; OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION: US-08-471-033-35
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US-08-471-044-35
; Sequence 35, Application; Sequence 35, Application; Sequence 35, Application; Sequence 35, Applicant: Warren, Applicant: Warren, Applicant: Mullins, Applicant: Nye, Goi; Applicant: Nye, Goi;
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Best Local
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Value

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/471,044
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STRANDEDNESS: sin
TOPOLOGY: linear
MOLECULE TYPE: othe
DESCRIPTION: /des
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
TELECOMMUNICATION INFORMATION: TELEPHONE: 919-541-8582
                                                                FILING DATE: 25-MAR-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                 APPLICATION NUMBER: US 08/218,018 FILING DATE: 23-MAR-1994 PRIOR APPLICATION DATA:
                                                                                                                                                           APPLICATION NUMBER: 1
FILING DATE: 09-SEP-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                 FILING DATE: 06-JUN-
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
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ADDRESSEE: CIBA-GEIGY Corporation
STREET: 7 Skyline Drive
                                                                                                                                                                                                       APPLICATION NUMBER: US 08/463,483 FILING DATE: 05-JUN-1995 PRIOR APPLICATION DATA:
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                            NAME: Pace, Gary M.
REGISTRATION NUMBER: 40,403
REFERENCE/DOCKET NUMBER: CGC
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CITY: Hawthorne
STATE: NY
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L4; Conservative
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T: Estruch, Juan J
INVENTION: No. 5840868el Pesticidal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GACTACTACATCAGC
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Mullins, Martha A
Nye, Gordon J
Carr, Brian
Desai, Naliri
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Kostichka, N.
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4: 800
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09-SEP-1994
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93.38;
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= "Synthetic DNA"
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encoding VIP1A(a) with the Bacil
contained in pCIB5526"
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Kristy
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                             1695/CIP3/DIV6
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; NAME/KEY: CDS; LOCATION: 9..2564; OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION: US-08-471-044-35
Дb
                    УV
                                        Best Loc
Matches
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                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 2576 base pair
                                                                                                                                                   MOLECULE TYPE:
DESCRIPTION:
HYPOTHETICAL:
FEATURE:
                                                   Local
                                                                                                                                                                                                  TOPOLOGY:
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1944 GACTACTACATCAGC 1958
                                                    Similar
        GACTACTACATGAGC 15
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                                        89.3%; illarity 93.3%; Conservative
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/desc = "Synthetic
                                                                                              /note= "Maize optimized sequence encoding VIP1A(a) with the Bacili contained in pCIB5526"
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                                                    Score 13.4;
Pred. No. 85;
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Search completed: June 26, 2003, 04:52:15 Job time: 24.2 secs

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1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*

2: /cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*

3: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*

5: /cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*

6: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*

7: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*

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11: /cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*

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13: /cgn2_6/ptodata/1/pubpna/US10_PUBCOMB.seq:*

14: /cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
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US-09-935-943-3

US-09-935-943-10

US-09-935-943-10

US-09-918-995-9782

US-09-918-995-9782

US-10-029-180-63

US-09-933-267A-1

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    sequence 25, Appl sequence 10, Appl sequence 175, Appl sequence 9782, Appl sequence 63, Appl sequence 63, Appl sequence 1, Appli sequence 1, Appli sequence 1, Appli sequence 23136, A sequence 6421, Apsequence 6624, Apsequence 1862, Apsequence 1862, Apsequence 1862, Apsequence 282, Appsequence 1769, Apsequence 1769, Appexequence 1769, Appexe
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sequence 850, App	equence	e ib, A	0 582,	3044,	e 15054,	7007	7002	0.42	Sequence 108/, Ap	e 3481/	Sequence 35180, A	Sequence 15/2, Ap	ce 541	Sequence 238, App	657,	ce 224	equenc	Sequence 281, App	ce 152,	1584,	e 2012,	ന	1847,	19, Ap	

ALIGNMENTS

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US-09-988-115A-52; Sequence 52, App.; Publication No.
                                                                                                                                 ; TYPE: DNA
; ORGANISM: Bovin
US-09-988-115A-52
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CURRENT FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: US 60/311,625
PRIOR FILING DATE: 2001-08-09
PRIOR APPLICATION NUMBER: US 60/256,458
PRIOR FILING DATE: 2000-12-20
PRIOR APPLICATION NUMBER: US 09/714,185
PRIOR APPLICATION NUMBER: US 60/166,410
PRIOR APPLICATION NUMBER: US 60/166,410
PRIOR FILING DATE: 1999-11-19
                                                                                                                                                                                  NUMBER OF SEQ ID NO
SOFTWARE: FastSEQ f
SEQ ID NO 52
LENGTH: 393
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APPLICANT: Goldsby, Richard A.
APPLICANT: Ferguson, Stacy E.
APPLICANT: Kuroiwa, Yoshima
APPLICANT: Tomizuka, Kazuma
APPLICANT: Ishida, Isao
                                                                Query Match
Best Local S
Matches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Ishida, Isao
TITLE OF INVENTION: Expression of Xenogenous (Human)
TITLE OF INVENTION: Immunoglobulins in Cloned, Transgenic Ungulates
FILE REFERENCE: 50195/008003
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TYPE: DY
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US20030037347A1
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Pred. No. 38;
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RESULT

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RESULT 3
US-09-938-842A-3563/c
; Sequence 3563, Application
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US20020160378A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; FEATURE:
; OTHER INFORMATION:
US-09-917-800A-1490
         TITLE OF INVENTION: STRESS-REGULATED GENES OF PL;
TITLE OF INVENTION: SAME, AND METHODS OF USE;
FILE REFERENCE: SCRIP1300-3
CURRENT APPLICATION NUMBER: US/09/938,842A
CURRENT FILING DATE: 2001-08-24
PRIOR APPLICATION NUMBER: US 60/227,866
PRIOR FILING DATE: 2000-08-24
PRIOR FILING DATE: 2001-08-24
PRIOR APPLICATION NUMBER: US 60/264,647
PRIOR APPLICATION NUMBER: US 60/300,111
PRIOR APPLICATION NUMBER: US 60/300,111
PRIOR FILING DATE: 2001-06-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Molecular Toxicology Notice of the Reference: 44921-5038-US CURRENT APPLICATION NUMBER: US/09/917,8007 CURRENT FILING DATE: 2001-07-31 PRIOR APPLICATION NUMBER: US 60/222,040 PRIOR FILING DATE: 2000-07-31 PRIOR APPLICATION NUMBER: US 60/222,880 PRIOR APPLICATION NUMBER: US 60/222,880 PRIOR APPLICATION NUMBER: US 60/290,029 PRIOR FILING DATE: 2000-11-02 PRIOR FILING DATE: 2001-05-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/295,798
PRIOR FILING DATE: 2001-06-06
PRIOR APPLICATION NUMBER: US 60/297,457
PRIOR FILING DATE: 2001-06-13
PRIOR APPLICATION NUMBER: US 60/298,884
PRIOR FILING DATE: 2001-06-19
PRIOR APPLICATION NUMBER: US 60/303,459
PRIOR FILING DATE: 2001-07-09
NUMBER OF SEQ ID NOS: 1740
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Matches 1
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ORGANISM: Rattus
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RESULT 5
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NAME/KEY: CDS;
LOCATION: (1)...(
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; TYPE: DNA
; ORGANISM: Arabidopsis thaliana
US-09-938-842A-3563
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US-09-935-943-3
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PRIOR APPLICATION NUMBER: PCT/EP01/08910;
PRIOR FILING DATE: 2001-08-01;
PRIOR APPLICATION NUMBER: US 60/222,779;
PRIOR FILING DATE: 2000-08-03;
NUMBER OF SEC. TO THE POINT NUMBER: US 60/222,779;
                                      PRIOR FILING DATE:
NUMBER OF SEQ ID NO
SOFTWARE: PatentIn
SEQ ID NO 25
LENGTH: 3236
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Best Local S
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APPLICANT: Levin, Joshua Z.
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GENERAL INFORMATION:
APPLICANT: Levin, Joshua Z.
APPLICANT: Wegrich, Lynette M.
APPLICANT: Budziszewski, Gregory J.
TITLE OF INVENTION: Herbicide Target Genes
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                                                                                                            FILE REFERENCE: PB/5-31509A
CURRENT APPLICATION NUMBER: US/09/935,943
CURRENT FILING DATE: 2001-08-23
PRIOR APPLICATION NUMBER: PCT/EP01/08910
PRIOR FILING DATE: 2001-08-01
PRIOR APPLICATION NUMBER: US 60/222,779
                                                                                                                                                                                                 APPLICANT: Wegrich, Lynette M.
APPLICANT: Budziszewski, Gregory J.
TITLE OF INVENTION: Herbicide Target Genes
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TYPE: DNA
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TYPE: DNA
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FEATURE:
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RESULT 6
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APPLICANT: Budziszewski, Gregory J.
TITLE OF INVENTION: Herbicide Target Genes an
FILE REFERENCE: PB/5-31509A
CURRENT APPLICATION NUMBER: US/09/935,943
CURRENT FILING DATE: 2001-08-23
PRIOR APPLICATION NUMBER: PCT/EP01/08910
PRIOR FILING DATE: 2001-08-01
PRIOR APPLICATION NUMBER: US 60/222,779
PRIOR FILING DATE: 2000-08-03
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TYPE: DNA
ORGANISM: Arabidopsis
09-935-943-10
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APPLICANT: Hoffman, Neil
APPLICANT: Hurban, Patrick
TITLE OF INVENTION: Expressed Sequences
TITLE OF INVENTION: thaliana
FILE REFERENCE: 2031US (PARA-020PRV)
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Matthew, Abraham
Ledford, Brooke
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Slader, Ted
Davis, Keith R.
Allen, Keith
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Hamilton, Carol
Price, Jennifer
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OR APPLICATION NUMBER: PCT/US01/00666

OR APPLICATION NUMBER: PCT/US01/00666

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Rank, David R.
Hanzel, David K.
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DATE: 2001-01-26

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NUMBER: US 0
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JIMBER: US 60/207,456
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APPLICANT: Hyseq, Inc.
FITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUE
FILE REFERENCE: 20411-756
FILE REFERENCE: 20411-756
CURRENT APPLICATION NUMBER: US/09/918,995
CURRENT FILING DATE: 2001-07-30
PRIOR APPLICATION NUMBER: US/09/235,076
PRIOR FILING DATE: 1999-01-20
NUMBER OF SEQ ID NOS: 38054
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 9782
LENGTH: 470
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US-09-918-995-9782
; Sequence 9782, Application US/09918995
; Publication No. US20030073623A1
; GENERAL INFORMATION:
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ORGANISM: Homo sapiens
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NAME/KEY: misc_feature
LOCATION: (1)...(470)
OTHER INFORMATION: n =
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184 GACTCCTACATGAGC 198
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104 GACTACAACATGAGC 118
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Silva, Jeffry C
Trueheart, Josh
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Pred. No. 3.2e+02;
); Mismatches 1;
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N ADULT LIVER, SIGNAL = 34
N BRAIN, SIGNAL = 44
N HEART, SIGNAL = 1.2e+02
N BT474, SIGNAL = 1.3e+02
N HBL100, SIGNAL = 1.3e+02
N FETAL LIVER, SIGNAL = 46
N LUNG, SIGNAL = 68
N HELA, SIGNAL = 77
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RESULT 11
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; OTHER INFORMATION:
US-10-029-180-63
                                                                                                  ; NAME/KEY: misc_feat; LOCATION: (1571); OTHER INFORMATION: US-09-925-301-93
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TITLE OF INVENTION: No. US20020182708A1e1
FILE REFERENCE: MIC-004
CURRENT APPLICATION NUMBER: US/10/029,180
CURRENT FILING DATE: 2001-12-22
PRIOR APPLICATION NUMBER: US 60/257,431
PRIOR FILING DATE: 2000-12-22
NUMBER OF SEQ ID NOS: 138
                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-08
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TYPE: DNA
ORGANISM: Artifi
                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/124,270 PRIOR FILING DATE: 1999-03-12
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                                                                                                                                                  NAME/KEY: misc_feat
LOCATION: (1546)
OTHER INFORMATION:
                                                                                                                                                                                       NAME/KEY: misc_feature
LOCATION: (1530)
OTHER INFORMATION: n equals
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OTHER INFORMATION: n equals
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LOCATION: (1497)
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              1 GACTACTACATGAGC 15
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Pred. No. 3.4e+02;
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RESULT
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; ORGANISM: human
US-09-933-267A-1
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; ORGANISM: C
US-09-738-626-1
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                                                                           CURRENT APPLICATION NUMBER: US/09/738,620
CURRENT FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: JP 99/377484
PRIOR FILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: JP 00/159162
PRIOR FILING DATE: 2000-04-07
PRIOR APPLICATION NUMBER: JP 00/280988
PRIOR APPLICATION NUMBER: JP 00/280988
PRIOR FILING DATE: 2000-08-03
NUMBER OF SEQ ID NOS: 7059
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TYPE: DN
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)R APPLICATION NUMBER: 09/692414
)R FILING DATE: 2000-10-20
)R APPLICATION NUMBER: 09/768184
)R FILING DATE: 2001-01-24

OR FILING DATE: 2001-01-24
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APPLICATION NUMBER: 09/826314
FILING DATE: 2001-04-05
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HAYASHI, MIKINO
OCHIAI, KEIKO
YOKOI, HARUHIKO
METSHI, NAOKO
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SENOH, AKI
IKEDA, MAS
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TITLE OF INVENTION: HUMAN GENOME-DERIVED OF INVENTION: GENE EXPRESSION ANALY
FILE REFERENCE: Aeomica-X-1
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
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N NUMBER: PCT/US01/00661
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2000-08-03
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: EXPRESSED IN PLACENTA, SIGNAL = 1.7
: EXPRESSED IN BONE MARROW, SIGNAL = 1.7
: EXPRESSED IN ADULT LIVER, SIGNAL = 1.7
: EXPRESSED IN HEART, SIGNAL = 1.3
: EXPRESSED IN FETAL LIVER, SIGNAL =
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UMBER: US 60/207,456
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FILE DE INVENTION: GENE LAPRESSION ANALISA
FILE REFERENCE: AGOMICA-X-1
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
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PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2000-09-27
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PRIOR APPLICATION NUMBER: US 09/608,408
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ISM Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

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Brunstein,A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and Simpson,A.J.

Shotgun sequencing of the human transcriptome with ORF expressed sequence tags

Contact: Simpson A.J.G.

Laboratory of Cancer Genetics

Ludwig Institute for Cancer Research

Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,

Brazil

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REFERENCE
AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Eco RI site shown at the beginning of the sequence.

Tissue Procurement: Louis M. Staudt, M.D., Ph.D.

cDNA Library Preparation: M.B. Soares Lab

cDNA Sequencing by: M.B. Soares Lab

Clone distribution: MGC clone distribution information car

found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

Seq primer: M13 Forward.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homc.

1 (bases 1 to 487)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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UI-HF-BK0-abo-a-12-0-UI.r1 NIH_MGC_
IMAGE:3056926 5', mRNA sequence.
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This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=&t2=MR3-UM0065-030
300-107-a09&t3=2000-03-03&t4=1)
Seq primer: puc 18 forward
High quality sequence stop: 308.
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/db_xref="taxon:9606"
/clone="IMAGE:3056926"
/clone_lib="NIH_MGC_36"
/tissue_type="lymph"
/cell_type="germinal center B cells"
/cell_line="MGC85"
/lab_host="DH10B (LTI)"
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/note="Organ: uterus; Vector: puc18; Site_1: SmaI; Site_2: SmaI; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."

89 c 81 g 73 t
/lab_host="DH10B (LTI)"

/lab_host="DH10B (LTI)"

/note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: Eco
/note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: PT7T3-Pac; Site_2: PT7T3-Pac; Site_1: NotI; Site_2: PT7T3-Pac; Site_2: PT7T3-Pac
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/db_xref="taxon:9606"
/clone_lib="UM0065"
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Pred. No. 9.8e+02;
Mismatches 0;
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cDNA Library Preparation: Ling Hong/Rubin Laboratory

cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can be
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AGENCOURT_8501500 NIH_MGC_113
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Email: cgapbs-r@mail.nih.gov
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                                                                                                                                                                                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="iMAGE:4763274"
/clone=lib="NIH_MGC_48"
/clone_lib="NIH_MGC_48"
/clone_lib="NIH_MGC_48"
/tissue_type="primary B-cells from tonsils (cell line)"
/tissue_type="primary B-cells from tonsils (cell line)"
/lab_host="DH10B (phage-resistant)"
/note="Organ: B-cells; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the
following 5' adaptor: GGCACGAG(G). Size-selected >500bp
for average insert size 1.8kb. Library constructed by Ling
Hong in the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
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Institutes of Health, Mammalian Gene Collection (MGC)
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Pred. No. 1.3e+03;
Mismatches 0;
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BQ710859
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602733346F1 NIH_MGC_,
mRNA sequence.
BG760090
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Tissue Procurement: Dr. Mark Watson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cDNA Library Preparation: Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Agencourt Bioscience Corporation Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLCM2521 row: i column: 14
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National Institutes of Health, Mammalian
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National Institutes of Health, Mammalian
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
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Mammalia; Eutheria; Primates;
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/note="Organ: spleen; Vector: pOTB7; Site_1: XhoI; Site_2:
EcoRI; cDNA made by oligo-dT priming. Directionally cloned
into EcoRI/XhoI sites using the following 5' adaptor:
GGCACGAG(G). Library constructed by Ling Hong in the
laboratory of Gerald M. Rubin (University of California,
Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and
Superscript II RT (Life Technologies). Note: this is a
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/db_xref="taxon:9606"
/clone="IMAGE:6303061"
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Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the
following 5' adaptor: GGCACGAG(G). Size-selected >500bp
for average insert size 1.8kb. Library constructed by Ling
Hong in the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
/organism="Homo sapiens"
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Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the
following 5' adaptor: GGCACGAG(G). Size-selected >500bp
for average insert size 1.8kb. Library constructed by Ling
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Unpublished (1999)
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15; Conservative
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URL:http://genome.gsc.riken.go.jp/
Sasaki,N., Izawa,M., Watahiki,M., Ozawa,
Matsuura,S., Carninci,P., Muramatsu,M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Yoshihide Hayashizaki
Laboratory for Genome Exploration Research Group, RIKEN Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-1
Tel: 81-45-503-9222
Fax: 81-45-503-9216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       house mouse
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                                                                                                                                                                         details.
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Note: this is a NIH_MGC Library."

313 c 330 g 236 t 1 others
/organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="4931436E24"
/clone_lib="RIKEN full-length enriched,
                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GI:6260848
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.1.7e+03;
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Okazaki,Y. and
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Hayashizaki
         male
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Fungal Ge
21110325
l Similar:
14; Cons
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The rpl16a Gene for Ribosomal Protein L16A Identified from Expressed Sequence Tags Is Differentially Expressed during Development of Aspergillus nidulans
Fungal Genet. Biol. 31 (2), 69-78 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Fungi; Ascomycota; Pezizomycotina; Eurotiomycetes; Eurotiales; Trichocomaceae; Emericella.

1 (bases 1 to 250)
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larity 100.0%;
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of Biological Sciences
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Emericella nidulans FGSC4 early sexual Emericella nidulans
ne esd0652, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          was cloned into the XhoI and BamHI sites. Vector: a modified pBluescript KS(+) after bulk excision from Lambda FLC I. Cloning sites, 5' end: SalI; 3' end: BamHI."

37 c 46 g 77 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     prepared and sequenced in Mouse Genome Encyclopedia
Project of Genome Exploration Research Group in Riken
Genomic Sciences Center and Genome Science Laboratory
RIKEN. Division of Experimental Animal Research in Rik
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    contributed to prepare mouse tissues. 1st strand primed with a primer [5' GAGAGAGAAGAAGCTCTTTTTTTTTTTTTTTTTTVN 3'],
                                                                                          /clone_lib="Emericella nidulans FGSC4 early
/dev_stage="early sexual"
/note="3'-directed cDNA clones; single-pass
                                                                                                                                       /organism="Emericella nidulans"
/strain="FGSC4"
/db_xref="taxon:162425"
/clone="esd0652"
                                                                                                                                                                                                                      Location/Qualifiers
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/tissue_type="testis"
/dev_stage="adult"
/lab_host="DH10B"
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              93.3%;
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Score 14; DB; Pred. No. 3e+0; Mismatches
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Pred. No. 2.9e+03;
0; Mismatches 0;
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Contact: Yoshihide Hayashizaki
Laboratory for Conoma Fynloration Basearch Group Divin Conomic
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AV272672.1
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Mammalia; E
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AV272672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Transcriptional sequencing: A method for DNA sequencing using RNA polymerase. Proc. Natl. Acad. Sci. U.S.A. 95 (7), 3455-3460 (1998) Itoh, M., Kitsunai, T., Akiyama, J., Shibata, K., Izawa, M., Kawai, J., Tomaru, Y., Carninci, P., Shibata, Y., Ozawa, Y., Muramatsu, M., Okazaki, Y. and Hayashizaki, Y. Automated filtration-based high-throughput plasmid preparation system. Genome Res. 9 (5), 463-470 (1999)

Carninci, P. and Hayashizaki, Y. High-efficiency full-length cDNA cloning. Methods Enzymol. 303, 19-44 (1999)

Please visit our web site (http://genome.rtc.riken.go.jp) for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Laboratory for Genome Exploration Research Group, RIKEN Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-Tel: 81-45-503-9222
Fax: 81-45-503-9216
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URL:http://genome.gsc.riken.go.jp/
Sasaki,N., Izawa,M., Watahiki,M., Ozawa,K., Tanaka,T.,
Matsuura,S., Carninci,P., Muramatsu,M., Okazaki,Y. and
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malia; Eutheria;
(bases 1 to 251)
                                                                                                                                                                                                                                                                                                                        /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="4931431F16"
/clone_lib="RIKEN full-le
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                                               /note="Site_1: Sall; Site_2: BamHI; cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken contributed to prepare mouse tissues. 1st strand cDNA was
Location/Qualifiers
                                                                                                                                                                                                             /tissue_type="testis"
/dev_stage="adult"
/lab_host="DH10B"
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Kagawa, I., Kai
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C50318 Yuji Kohara unpuk
C50318 Yuji Kohara unpuk
Caenorhabditis elegans c
C50318
C50318.1 GI:2387571
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C50628 Yuji Kohara unpublished cDNA:Strain N2 hermaphrodite embryo
Caenorhabditis elegans cDNA clone yk497b1 5', mRNA sequence.
C50628
C50628.1 GI:2387881
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Caenorhabditis elegans
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Biology Lab.

al Institute of Genetics

111, Mishima, Shizuoka 411, J

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/sex="hermaphrodite"
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tidae; Peloderinae; Caenorhabditis.
s 1 to 360)
                   377 bp mRNA linear EST 18-OCT-1999
Juji Kohara unpublished cDNA:Strain N2 hermaphrodite embryo bditis elegans cDNA clone yk480c3 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                kohara@lab.nig.ac.jp.
Location/Qualifiers
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/db_xref="taxon:6239"
/clone="yk497b1"
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Yata 1111, Misnim,
Tel: 81-559-81-6854
Fax: 81-559-81-6855
Email: ykohara@lab.nig.ac.jp.
Location/Qualifiers
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,M., Miyata,A. and Nishigaki,A.

Expression map of the C.elegans genome
Unpublished (1996)
Contact: Yuji Kohara
Genome Biology Lab.
                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 405)

Dias Neto, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R., Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F., Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H., Brunstein, A., deOliveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and Simpson, A.J.

Shotgun sequencing of the human transcriptome with ORF expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BF919229 405 bp mRNA linear CMO-NT0131-281000-641-g10 NT0131 Homo sapiens cDNA, BF919229 BF919229.1 GI:12315117 EST.
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Caenorhabditis elegans
Caenorhabditis elegans
Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL (http://www.ludwig.org.br/scripts/gethtml2.pl?tl=CMO&t2=CMO-NTOl31 281000-641-g10&t3=2000-10-28&t4=1)
                                                                                                                                           Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 and
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                                                                                        Tel: +55-11-2704922
Fax: +55-11-2707001
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/clone_lib="Yuji Kohara unpublished cDNA:Strain N2
/exmaphrodite embryo"
/sex="hermaphrodite"
/dev_stage="embryo"
76 c 86 g 102 t 1 others
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/db_xref="taxon:6239"
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Other_GSSs: CIT-HSP-2303C19.TF
Contact: Mark Adams
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Clones are available fro
end search page:
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/clone="2303C19"
/clone_lib="CIT-HSP"
/sex="Male"
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www.tigr.org/tdb/humgen/bac_end_search/bac_end_search.html.
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                                       /cell_type="Sperm"
/note="Vector: pBe
HindIII"
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/db_xref="taxon:9606"
/clone_lib="NT0131"
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andom BAC End Sequence Database for Sequence Database fo
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for Genomic Research
Center Dr., Rockville,
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Proc. Natl. Acad. Sci. U.S.A. 97
20202663
Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Resea
Rua Prof. Antonio Prudente 109, 4
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MR3-CI0185-
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Eukaryota; Me
Mammalia; Eut
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Similarity 100.0%;
14; Conservative
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1 (bases 1 to 417)

1 (bases 1 to 417)

Dias Neto, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R., Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F., Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H., Brunstein, A., deOliveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and
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Shotgun sequencing of
                                                                                                                        Similarity 100 14; Conservative
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061200-002-b11&t3=2000-12-06&t4=1)
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Location/Qualifiers
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                                                                                                                                                                                          /note="Organ: colon_ins; Vector: pucl8; Site_1: Smal; Site_2: Smal; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."

109 c 143 g 80 t
                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens
/db_xref="taxon:9606"
/clone_lib="CI0185"
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Fax: 314 286
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: est@watson.wustl.edu
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further:
Trace considered overall poor quality
Seq primer: -40m13 fwd. ET from Amersham
High quality sequence stop: 1.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                  /dev_stage="20 week-post conception fetus"
/lab_host="DH10B (ampicillin resistant)"
/note="Organ: Liver and Spleen; Vector: pT7T3D (Pharmacia)
with a modified polylinker; Site_1: Pac I; Site_2: Eco RI;
This is a subtracted version of the original Soares fetal
liver spleen 1NFLS library. 1st strand cDNA was primed
with a Pac I - oligo(dT) primer [5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="GDB:1351880"
/db_xref="taxon:9606"
/clone="IMAGE:447623"
/clone_lib="Soares_fetal_liver_spleen_1NFLS_S1"
/sex="male"
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 to 420)
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15: em_ba:*

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Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Salonen, E.-M., Parren, P.W.H.I., Graus, Y.F., Fisicaro, P., Vapalahti, O., Kallio-Kokko, H., Burton, D.R.
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Salonen, E.-M., Parren, P.W.H.I., Graus, Y.F., Lunc Fisicaro, P., Vapalahti, O., Kallio-Kokko, H., Vahe Burton, D.R.
Direct Submission
Submitted (09-OCT-1997) Immunology, The Scripps 10550 N. Torrey Pines Road, La Jolla, CA 92037, Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                  Human recombinant Puumala virus antibodies: other hantaviruses and use in diagnostics Unpublished
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Mammalia; Eutheria; Primates; Catarrhini; Homini
CE 1 (bases 1 to 717)

RS Merchant, A.M., Zhu, Z., Yuan, J.Q., Goddard, A., Ada
Presta, L.G. and Carter, P.
An efficient route to human bispecific IgG
Nat. Biotechnol. 16 (7), 677-681 (1998)
E 98325681
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NEDO human cDNA
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Submission
ed (04-JUL-2002) Takao Isogai, FLJ Project(HRI Team);
Kamatari, Kisarazu, Chiba 292-0812, Japan
genomics@hri.co.jp, Tel:81-438-52-3975, Fax:81-438-5
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/protein_id="AAC98736.1"
/protein_id="AAC98736.1"
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IYHWLAWYQQKPGKAPKLLIYKASSLASGAPSRFSGSGSGTDFTLTISSLQPDDFATY
YCQQYSNYPLTFGGGTKLEIK"
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Davies, C.J., Davis, C., English, C., Fondon, T., Franklin, T.L.,
Garner, H.R., Gordon, M., Gotway, G., Grant, O., Hanher, L., Harris, J.,
Hinson, S., McFarland, J., Megarity, C., Narayanaswamy, U., Newton, J.,
O'Brien, K., Oliver, T., Patel, P., Probst, S., Rayner, S.,
Schageman, J., Schilling, P., Schultz, R., Syed, M., Valenzuela, D.,
Ward, T. and Wilson, R.
Direct Submission
Liversity of Texas Southwestern Medical Center at
Development, University of Texas Southwestern Medical Center at
Dallas, 5323 Harry Hines Blvd, Dallas, TX 75235-8591, USA
On Jun 19, 1997 this sequence version replaced gi:1737188.
Location/Qualifiers
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15; Conser
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Mammalia; E
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Human chromosome 11 146h12 cosmid,
U73633
U73633.1 GI:2199506
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malia; Eutheria; Primates;
(bases 1 to 42845)
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1. .2030
                                                                                                                                                                       /clone_lib="library prepared from flochromosome 11 derived from Chinese has monochromosomal somatic cell hybrid, 2218. . 2488
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AP000590.
Homo sapiens DNA, clone: CMB9-52H4
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15; Cons
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milarity 100.0%;
Conservative 0
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/rpt_type=dispersed
complement (24053. .2:
/rpt_family="ALU"
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26266. .26540
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complement(28625. .28911)
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/rpt_family="MER46"
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a 11179 c 11246 g 10800 t
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39388. 39646
/rpt_family="ALU"
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35218
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Pred. No. 1.7e+03;
Mismatches 0;
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Direct Submission
Submitted (12-OCT-1999) Masahira Hattori, The Institute of Physical Research (RIKEN), Genomic Sciences Center (GSC);
and Chemical Research (RIKEN), Genomic Sciences Center (GSC);
1-7-22 Suehiro-chou, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan (E-mail:hattori@gsc.riken.go.jp, URL:http://hgp.gsc.riken.go.jp/,
Tel:81-45-503-9111, Fax:81-45-503-9170)
On Jul 17, 2001 this sequence version replaced gi:10880434.
Location/Qualifiers
1. .114287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence.
AC018902
AC018902.8
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Bloom, S.,
                                      Pate,D. and Hood, L.

Direct Submission
Submitted (09-AUG-2001) Multimegabase Sequencing Center, Institute
Submitted Siology, 4225 Roosevelt Way NE, Suite 200, Seattle, WA
                                                                                                                                                      and Hood, L.

Direct Submission

Direct Submission

Submitted (22-DEC-1999) Multimegabase Sequencing of Washington, PO BOX 357730, Seattle, WA 98195, 3 (bases 1 to 157226)

Rowen, L., Madan, A., Qin, S., Baradarani, L., Birdit Burke, J., Dors, M., Fleetwood, P., Kaur, A., Madan, A. Pate, D. and Hood, L.
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2 (bases 1 to 114287)

Hattori,M., Ishii,K., Toyoda,A., Taylor,T.D., Fujiyama,A., Yada,T., Totoki,Y., Watanabe,H. & Fujiyama, Yada,T., Yada,T.,
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                       On Aug 9, 2001 this
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1 (bases 1 to 157226)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequencing of human chromosome 15 D15S146-D15S117 region Unpublished
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Eutheria;
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Dors, M., Dickhoff, R., Fleetwood, P., Harrison,
Kaur, A., Madan, A., Owen, M.P., Ratcliffe, A., S
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/db_xref="taxon:9606"
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Fujiyama
                           Submitted (17-DEC-1999) Masahira Hattori, The Instit and Chemical Research (RIKEN), Genomic Sciences Cent Kitasato Univ., 1-15-1 Kitasato, Sagamihara, Kanagaw Japan (E-mail:hattori@gsc.riken.go.jp, URL:http://hgp.gsc.riken.go.jp/, Tel:81-42-778-9923,
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               URL:http://hgp.gsc.riken.go.jp/, Fax:81-42-778-9924)
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HTGS_PHASE1; HTGS_DRAFT.
sapiens DNA, clone:RP11-677D19.
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lencing vector: pUC18; L08752
listry: Dye-terminator Big Dye; 90% of reads
mistry: Dye-primer Big Dye; 10% of reads
mbly program: Phrap; version 0.990399
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/note="Data from overlapping BACs RP11-430B1, RP11-23
and RP11-472P3 were added and the consensus sequence
determined from RP11-327C2 to the extent possible"
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NOTE: This i 61 contigs. order in thi contigs are contigs are are unknown. as soon as i preserved
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Contact: hattori@gsc.riken.go.jp/
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Contact: hattori@gsc.riken.go.jp/
Contact: hattori@gsc.riken.go.jp/
Conter project Information
Center project name: HumDraftll
Center clone name: RP11-677D19

Center clone name: RP11-677D19

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AP001271
AP001271.4 GI:15320469
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Published Only in Database (2000)
2 (bases 1 to 210402)
Hattori, M., Ishii, K., Toyoda, A., Taylor, T.D., Hong-Seog, P., Fujiyama, A., Yada, T., Totoki, Y., Watanabe, H. and Sakaki, Y. Direct Submission
Submitted (23-FEB-2000) Masahira Hattori, The Institute of Physical and Chemical Research (RIKEN), Genomic Sciences Center (GSC); 1-7-22 Suehiro-chou, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan (E-mail:hattori@gsc.riken.go.jp, URL:http://hgp.gsc.riken.go.jp/, Tel:81-45-503-9111, Fax:81-45-503-9170)
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Similarity 100.0%;
15; Conservative 0;
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Eukaryota; Metazoa; Chordata; Crania
Mammalia; Eutheria; Primates; Catar:
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                                                                 /organism="Homo sapiens"
/db_xref="taxon:9606"
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15; Conser
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Direct Submission
Submitted (17-SEP-1999) Multimegabase Sequencing Center, University of Washington, PO BOX 357730, Seattle, WA 98195, USA
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                                                                                                                                                                                                                                                                                                                              web site: http://chroma.mbt.washington.edu/msg_www Contact: leerowen@systemsbiology.org
------- Summary Statistics
Sequencing vector: pUC18; L08752
Chemistry: Dye-terminator Big Dye; 90% of reads
Chemistry: Dye-primer Big Dye; 10% of reads
Assembly program: Phrap; version 0.990399
                                                                                                                                                                                                                                                                                                                                                                                                                       Center: Multimegabase Sequencing Center Center code: UWMSC Web site: http://chroma.mbt.washington.contact: leerowen@systemsbiology.org
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submission
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/clone_lib="RPCI human BAC library 11"
/note="Data from overlapping BACs CTD-2184D3, RP11-327C2, and RP11-337B11 were added and the consensus sequence was determined from RP11-430B1 to the extent possible"
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'note="overlap with CTD-2184D3 AC023906"
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Pred. No. le+03;
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Direct Submission

L Submitted (26-JUL-2001) Gouzy J., Submitted on behalf of the MELILO EU Consortium

MELILO EU Consortium:

Laboratoire de Biologie Moleculaire des Relations

Plantes-Microorganismes, UMR215-CNRS-INRA, BP27, F-31326 Castanet, France, Laboratoire de Genetique et Developpement UMR6061-CNRS, France, GATC GmbH, Fritz-Arnold-str. 23, D-78467 Konstanz, Germany, Universitaet Bielefeld, Biologie IV (Genetik) Universitaetstr 25, D-33615 Bielefeld, Germany, Unite de Biochimie physiologique, Universite Catholique de Louvain, Place Croix du Sud 2, Bte 20, B-1348 Louvain-la-Neuve, Belgium, Unite de Microbiologie, Faculte des Sciences Agronomiques de Gembloux, Avenue Marechal Juin 6, B-5030 Gembloux, Belgium. E-mail:Jerome.Gouzy@toulouse.inra.fr http://sequence.toulouse.inra.fr/meliloti.html.

1. .333800
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21396507
11481430
2 (bases 1
Gouzy,J.
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15; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sinorhizobium meliloti.
Sinorhizobium meliloti
Sinorhizobium meliloti
Bacteria; Proteobacteria; alpha subdivision; Rhizobiaceae
Rhizobiaceae; Sinorhizobium.
1 (bases 1 to 333800)
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Sinorhizobium meliloti 1021 complete chromosome;
AL591792 AL591688
AL591792.1 GI:15075850
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/note="low quality data"

146876...146893

/note="low quality data"

165370...165440

/note="low quality data"

177221...214415

/note="overlap with RP11-:

a 47930 c 46801 g 57867
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/note="low quality data"
119015. .119515
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/note="sequence
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57867 t
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9.9e+02;
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1305. . 167
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2459. 4297
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /strain="1021"
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Gene name confidence : hypothetical
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Gene name confidence : hypothetical
predicted by Codon_usage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /transl_table=11
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EPFSLLLTDIVMPEMDGIELARRATELDPDLKVMFITGFAAVALNPDSKAPKDAKVLS
KPFHLRDLVNEVNKMLAA"
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complement(233...1114)
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                                                                                                                                                                                                                     biosynthesis; isoleucine/v/note="Product confidence
Gene name confidence : pro
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                                                                                                                                                            predicted by
                                                                                                                                                                                                predicted by
                                                                                                                                                                               predicted
                                                                                                                                                                                                                                                                                                                                                                                                                         /gene="tRNA-VAL_GAC"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'evidence=not_experimental
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by FrameD"
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Homology
FrameD"
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"MPAYRSRTTTHGRNMAGARGLWRATGMKDSDFGKPIIAVVNSFT
                                                                                                                   _experimental
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/function="small molecule m
electron transport"
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Gene name confidence: puta
predicted by Codon_usage
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complement(5586. .6029)
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complement(5586. .6029)
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REPEAT SM-5
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GVDPAEFAALPIQYMDGLHNNWWNAPAETRHM"
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5027. .5084
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Gene name confidence : hypothetical
predicted by Codon_usage
predicted by FrameD"
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complement(6076. .6417)
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REPEAT SM-2
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5094. .5542
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                                                                                                                                                                                                                                            /evidence=not_experimental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Sm-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note="REP
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/db_xref="GI:15075855"
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disease
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Mammalia
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Khamlichi, A.A.
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larity 100.0%;
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/db_xref="taxon:9606"
/chromosome="2"
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/note="Product confidence : putative Gene name confidence : hypothetical
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/transl_table=11
/product="PUTATIVE CYTOCHROME C PROTEIN"
/protein_id="CAC47410.1"
/db_xref="GI:15075856"
/db_xref="SPTREMBL:Q92M25"
/translation="MTHASSRRIAAVLTCVTIGLSPRMASAEEDKLALGREIFLERSE
                                                             /product="immunoglobulin heavy
/note="gamma isotype"
233 c 225 g 158 t
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/note="gamma isotype"
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/note="gamma isotype"
58. .427
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/tissue_type="bone marrow"
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             DB 9;
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tig of 154 by of unknown lig of 124 by of 105 by of 105 by of unknown lig of 993 by tig of 961 by of unknown tig of 961 by of unknown tig of 40 by of unknown tig of 65 bp of unknown tig of 131 by of unknown tig of 131 by of unknown tig of 131 by of unknown tig of 46 bp of unknown tig of 131 by of unknown tig of 281 by tig of 281 by	* 17915 18752: contig of unknown length * 18753 18847: contig of 838 bp in gap of unknown length * 18753 18847: contig of 95 bp in length * 18848 19088: contig of 241 bp in gap of unknown length * 19089 19246: contig of 158 bp in gap of unknown length * 19247 19367: contig of 121 bp in gap of unknown length * 19368 19483: contig of 116 bp in gap of unknown length * 19484 19786: contig of 303 bp in gap of unknown length * 19787 20607: contig of 821 bp in gap of unknown length	11665 12907: contig of unk 12908 13764: contig of unk 13765 14507: contig of unk 13765 14507: contig of unk 14508 15239: contig of unk 15240 15636: contig of unk 15637 16504: contig of unk 16505 17834: contig of unk 17835 17914: contig of	cig of 193 bp of unknown laig of 78 bp of 65 bp of 639 bp of waknown laig of 885 bp of unknown laig of 263 bp of unknown laig of 197 bp of unknown laig of 197 bp of unknown laig of unknown laig of 197 bp of unknown laig of 427 bp of unknown laig of 427 bp

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Cenarchaeum symbiosum strain B histone H1 DNA binding protein (hc2), hypothetical protein, lysyl tRNA synthetase (syk), and hypothetical protein 01 genes, complete cds; 23S ribosomal RNA 16S ribosomal RNA genes, complete sequence; glutamate-1-semialdehyde aminotransferase (gsat) and hypothetical protein 02 genes, complete cds; tRNA-Tyr gene, complete sequence
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/db_xref="taxon:9606"
/chromosome="5"
/clone="CTB-163N20"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    J. Bacte
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9748430
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Feldman,
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Swanson,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             riol. 180 (19),
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                                                                                AAGVNEFSLSAPFDTTLPVHVELHDIGGQPAVDLAFAEDGRTLLLLAADGTLDFYSLA
GDAYDIGEASRTFQVPFEDAAGAVPGAFYQPPDGSSIIAAFDGRIDQYVVIPFEFVSY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TENAIRQYDLEGPYDIRGAGNAGQYDLDIPLHPGLLFLLTSGVHFSPDGTRMFVGEGI
SDAEDANANRDVNVNLWHRFDLSTPFDVLTAERVDTYEYSTGPAGDLEDLSLSPDGRR
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EFPFDVSSRTRADTLDIPQIDSPRHVAVSMPGNHLYITNSVFGEDDTIHSYGISNNDI
SSASYIGEEGIPEPVINGIDFSNNGRRMFLIGGNGFDYQVIHDYMLGTRYDISSRSLL
                                                                                                                                                                                                                                                       DVPSPSGIEFSDDGMRMFVTGIGTPGINIFTLSAPFDITLPKHSGSTNIGGLSVSDLAFANNGNSLTVLDVDGVLRVYALGDDYNVVTGTTQKFRITLDTTQGIPNSIYTSPDGLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                         PGVARFSLSDAEVLAVSGYAEPSLVFGRHAVPGAAGGTFPSQIGNATELVGSIPNPTL
DFGTTLTGAAFSADGTVVFLSDGPTGRVYPYSLNIPFDISSAAPGGFVIVPVGVSDIA
FSADGRNMLVADETGGIHRYLARSPYEIGTDFIKSSLGEFVETFSAAPRVQDLAGIAF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              YDLGAPYDISGPAYSGIFDAGGSIRDVAVGGGSMFILEGETDRVYEHRPGIYPVVSAL
DGPALVSAAADARVGAAEVLFDRAVDVGGIDPGGVRIVDAAGPLPGVVISDAVIPGED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GMDRSVHMYSLNTPFDLGGAEHAASFGVGDRVSDPLGTAFGNGGTKMLIADTTGFVHG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       YGDQSLHVPAAALAGTGGGPFVPDFSGGSLLASLYRHERPFQGEEMARTERSDRYALTVTAGGSQMHVGGAGGNITWYDLGTPHDITTGVRAGSDILPAYPSAGRNVVPSITGIAF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (07-AUG-1998) Genomics, Di., San Diego, CA 92121, USA Location/Qualifiers
                                                                                                                                                                     QYLLSEPFAVTTSVFLRTIPIDGGAEGIRFVDNGRGLFVPGADGTIQRHELIYPYGAS
TSLLETVRDGVTDGGPGENPAAGEIRLAGTFNASDNVQSPSGIEFSGDGTGMFVTGFG
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MPAPVMEQVTLDSREGTLRVRLDRTVDVDTVRPYKMWVEDSDGSQTTLANSTLLNAEN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TRLLVTGHGQTNAKLFHLNPPFDVGTAVFHDHGRFRPGGPASEIEASGISLSADGSRM
FLSDRGRGAISQYTLVAPFDVEFASDVSADGQLDVGAQDALPGGLAFSPGGTRLFMVG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SDMDGSPVGAGFAGG:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FGSSVYVGSDVAAPRGVYVAPGGSLMLVSDSADGTIHRYELASPYEPAGAANRGSFDV
SDMDGSPVGAGFAGGLHMYVAGNDTGRVYQYPAGTHQIQEAAAGPRLLSAVLDKDGTL
                                                                                                                                                                                                                                QFVAYDDRIDLYVLGSPNDISSTTEIIPYSLPRPDPPTGMDFTPDGRRMFLSTENGID
                                                                                                                                                                                                                                                                                                                  SNILLFRLDDAAAGKISGYTSPVFRTWSSPFLGTDGATRPHTLGFGDVRLADIYDASG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LYTLSSERVSSSEYTITRAQYWLPEPYDVTPPYHVPSFNASQGGNLADPYGMAFSPDG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SDDGMRLFAANRGDRIPMYQLDSPYDIGSASLEGTLFTGFQSGIAFSDDGTRMFAALL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RAAFDGTVDAGSVQPGMITIRDGHGSNTGIPLLLAGGAADSDVMTFVVPEKDRAEAAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'product="unknown"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'note="symbiont of the marine sponge Axinella mexicana"
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10625. .11434
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SVQKTS"
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PHPHHTRYEMFLDKGGKKISKSSGNVVTPQKWLRYGTPQSILLLLMYKRITGARELGLE
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VERSION
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                                                    TITLE
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Wray, P.
Direct Submitted
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Eukaryota;
Mammalia; E
                                                                                                                                                                       complete
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AL357123.
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Human DNA
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Similarity 100.0%;
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NWPYDVPESGGLVDEEHTVSIPYNNLEGSLEALRRAGGDLACVIVEPMLGGGGCIPAE
PDYLRGIQEFVHSKGALFILDEIVTGFRFDFGCAYKKMGLDPDVVALGKIVGGGFPIG
VVCGKDEVMCISDTGAHARTERAYIGGGTFSANPATMTAGAAALGALRERRGTLYPRI
NSMGDDARARLSRIFDGRVAVTGRGSLFMTHFTPDGARRISSAADAAACDVHLLHRYH
LDMITRDGIFFLPGKLGAISAAHSRADLGAMYSASERFAGGL"
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EKHPGRLGAWLPLLGQYPPGLVQKCISSQKLAEVLVQKWLEDHMFAGESDAAEKSKKI
SGMLASPGKYYSHGRYISREECRGIGLKITDLEADQEFQDLTLSVSHAADILSQFTPI
NKIIANHLGNSVISKPST"
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LSRLFRENRTERVTKKLVEYGVIDGPSPGIERLIALAGNYADDMYSAERTEVELDGAT
RGALSELAEMLGSAPEGGLQDVIYGVAKSHGVPPRDFFKALYRIILDASSGPRIGPFI
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/product="23S ribosomal RNA"
21771. .23243
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AIEASPRYAPAYFNRAVLLDRLGEHEDALPDLDKATRLDRDKANPRFYKGIVLGKMGR
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/transl_table=11
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                                                         Metazoa; Chordata;
Eutheria; Primates;
1 to 72354)
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 (08-MAR-2001)
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Pred. No
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Sanger
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Catarrhini;
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During sequ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Group. Further information can be found at http://www.sanger.ac.uk/HGP/Chrx
RP11-287E6 is from the library RPCI-11.1 constructed of Pieter de Jong. For further details see http://www.chori.org/bacpac/home.htm
VECTOR: pBACe3.6
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This sequence was finished as follows unless otherwise noted: all regions were either double-stranded or sequenced with an alternate chemistry or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by at least one plasmid subclone or more than one M13 subclone; and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           variation annotation may not be found i corresponding to the overlapping clone,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        On Mar 12, 2001 this sequence version replaced gi:13234914. During sequence assembly data is compared from overlapping clowhere differences are found these are annotated as variations together with a note of the overlapping clone name. Note that variation annotation may not be found in the sequence submissions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     database can be found at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    p://www.sanger.ac.uk/Projects/C_elegans/wormpep This si
generated from part of bacterial clone contigs of humo
omosome X, constructed by the Sanger Centre Chromosome
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/note="L2 rr

7760.
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/note="match: STS: Em:G60993
match: GSS: Em:AQ200790"
complement(916. .1482)
/note="match: STS: Em:HSC19C12
2401. .2612
                                                                /note="L1MC5
9537. .1000
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2762. .2804
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4774. .4900
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/chromosome="X"
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Location/Qualifiers
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/clone="RP11-287E6"
/clone_lib="RPCI-11.1"
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32756. .330
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35957. .:
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/note="MLT1F r
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33150. .3328:
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15763. .1640
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/note="Alusx repeat:
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ch 93.3%;
| Similarity 100.0%;
| 14; Conservative
                                                                    /note="LLM4 repeat: matches 1225..1319 of consensus"
141685.41978
/note="Alusg repeat: matches 1..306 of consensus"
41979.41992
/note="LIM4 repeat: matches 1213..1225 of consensus"
41998.42194
/note="Alusp repeat: matches 116..313 of consensus"
42199.43409
/note="LLM4 repeat: matches 116..313 of consensus"
42199.43525
/note="L2 repeat: matches 2399..2495 of consensus"
43430.43525
/note="L2 repeat: matches 2614..2750 of consensus"
43606.43736
/note="L2 repeat: matches 2614..2750 of consensus"
44810..46619
/note="L1 repeat: matches 1467..2749 of consensus"
46662.466419
/note="L1M4 repeat: matches 2..255 of consensus"
46662.46809
/note="L1M4 repeat: matches 4611..4760 of consensus"
46910.48311
/note="L1M4 repeat: matches 4755..6158 of consensus"
48376.48668
/note="L1ME2 repeat: matches 5386..5419 of consensus"
48669.49381
/note="PLIM52 repeat: matches 5386..5419 of consensus"
48669.49381
/note="L1M62 repeat: matches 5386..5419 of consensus"
48669.49381
/note="L1M62 repeat: matches 5419..6164 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="L1PA13 repeat: matches 523...
39502..40031
/note="L1M4 repeat: matches 4793...5
40379...41594
/note="L1M4 repeat: matches 2614...3
41599...41684
/note="L1M4 repeat: matches 1225...1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note="L1MB6 repeat: matches 5318.
37209. .37447
/note="L1PA15 repeat: matches 5918.
37475. .37531
/note="L1P repeat: matches 2395. .2/
37571. .39494
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41685.
    0;
    Score 14; DB 9; L
Pred. No. 4.3e+03;
); Mismatches 0;
                                     Length 72354;
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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388	ABA04555 ABK71369 ABA04563	A0455 K7135	AAX3239 ABK7136	ID
Anti-Mpl scFv 12B5	Human MPL #1 codin DNA encoding throm Human coding seque	Oligonucleotide #2	Ab3 variable heavy Thrombopoietin (TP	Description

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72	ABLIBS	N	2	9	Ü	44	
⊃	AAS9819	2	3735	စ	13.4	43	C
encoding	AAS9819	24	7	9	ω	42	
encoding	AAS9817	2	7	9	W	41	
\Box	AAZ8999	2	7	9	w	40	
ichia sp.	AAV6942	19	7	9	Û	39	
a mel	ABL0423	2	9	9	w	38	Ω
ch	AAC8643	21	9	9	w	37	Ω
starch	AAC8641	2	9	9	ω	ა 6	ဂ
Escherichia coli p	ABA8912	2	4	9	ω	S G	
Kidney cancer rela	ABL6854	24	25	9	£	34	a
Human cDNA differe	ABK8433	24	2	9	ω	ω ω	O
	AAH6679	22	74	9	w	32	O
Corynebacterium gl	AAF7230	2	9	9	w	31	O
	AAS7926	2	61	9	ω	30	
	AAH4356	2	3	9	ω	29	
Υt	AAT6826	1	ω	9.	w.	28	
0	AAX7579	20	W)	9	ω	27	
Human secreted pro	AAA3907	21	Ų.	9.	ω.	26	
0	AAH4356	22	w	9	w	25	
Œ	ABK3622	24	\mathbf{L}	9	w.	24	C
Porphorymonas ging	AAX9153	20	~	9	w	23	C
Human PTHrP monocl	AAA1393	21	N	9	ω.	22	
Human PTHrP monocl	AAA1393	21	N	9	ω ·	21	
Human PTHrP monocl	AAA1393	21	N	φ.	w.	20	
Human PTHrP monocl	AAA1393	21	N	<u>.</u>	ω	19	
Human PTHrP monocl	AAA1393	21	N	9	ω.	18	
_	AAH6863	22	\sim	9	·	17	
	AAH6862	22	\sim	9	ω.	16	
Human anti-Rh(D) c	AAH6861	22	\sim	œ •	·	5	
Corn tassel-derive	ABL7185	24	w		w •	14	O
Mouse spliced tran	ABN5782	24	65	9	ω.	<u>1</u> 3	
Cenarchaeum symbio	AΑ	21	42432	93.3	14	12	
3	AAA5519	21	57	ω.	14	11	
m	ABL5660	24	90		15	10	Q

ALIGNMENTS

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RESULT 1
AAX32399
ID AAX3
XX
AC AAX3
XX
DT 17-C
XX
ADE Ab3
XX
CW Mega
KW Mega
KW mega
KW myel
KW myel
KW war:
XX
PN WO9:
XX
PD 04-P
XX
PP 21-J
XX
PP 21-J
XX
PP 21-J
XX
PP A (GET
XX
PP Adam
XX
MPI
                                                                                                                                                                                        myelodysplasia;
neuromuscular; π
variable heavy c
                                                                                                                                                                    Homo
                        Adams
                                                                                                                                                                                                                                                                             Ab3
                                               (GETH
                                                                                                                                                                                                                                       Agonist antibod megakaryocyte;
                                                                                                                                                                                                                                                                                                                                                  AAX32399
                                                                                                                                                                                                                              bone marrow
                                                                                                                                                                                                                                                                                                   17-JUN-1999
                                                                                                                     04-MAR-1999.
                                                                                                                                            WO9910494-A2
                                                                      25-AUG-1997;
                                                                                              21-AUG-1998;
                                                                                                                                                                                                                                                                            variable heavy (VH) chain CDR1 encoding DNA.
                                                                                                                                                                  sapiens
  1999-204666/17.
                         CW,
                                                GENENTECH INC
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                         Carter PJ,
                                                                                                                                                                                                   lbody; thrombopoietin receptor; TPO-R; thrombopoietin; DIC; ce; platelet; immunological; hematopoietic; thrombocytopenia; hypoplasia; disseminated intravascular coagulation; anemia; sia; myelotoxic chemotherapy; leukaemia; tumour; MuSK; CDR; muscular dystrophy; complementarity determining region;
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                                                                                               98WO-US17364.
                                                                                                                                                                                            chain;
                                                                                                                                                                                                                                                                                                                                                   DNA;
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                                                                                                                                                                                            variable light chain;
                         Fendly BM,
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                          Gurney AL;
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P-PSDB;

AAY06699

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RESULT 2
ABK71361/c
ID ABK71361;
XX
AC ABK71361;
XX
AC ABK71361;
XX
DT 30-JUL-20
XX
DT 30-JUL-20
XX
DF Thrombopo
XX
Modified
KW TPO recep
KW TPO recep
KW ds.
XX
OS Synthetic
XX
PN WO2002330
XX
PD 25-APR-20
XX
PF 22-OCT-20
XX
PR 17-APR-20
PR 17-APR-20
PR 17-APR-20
XX
PR 112-SEP-20
XX
PA (CHUS) C
XX
PI Tsuchiya
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              megakaryocytes. They may also be able to stimulate megakaryocytes to increase platelet production. They can be used for treating immunological or hematopoietic disorders, especially thrombocytopenia. Thrombocytopenia -associated bone marrow hypoplasia (e.g. aplastic anemia C following chemotherapy or bone marrow transplant) may be effectively treated with the antibody compounds as well as disorders such as C disseminated intravascular coagulation (DIC), immune thrombocytopenia (HIV-induced and non HIV-induced), chronic idiopathic thrombocytopenia (congenital thrombocytopenia, thrombotic thrombocytopenia and C myelodysplasia. They can also be used in e.g. myelotoxic chemotherapy for C treatment of solid tumours or leukaemia, myeloablative chemotherapy for idiopathic aplastic anemia, congenital thrombocytopenia, and immune thrombocytopenia. The antibodies which bind to the Musk receptor can be used for improving neuromuscular function in a patient, e.q. in muscular function.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
Matches 1
                                                               20-OCT-2000;
17-APR-2001;
12-SEP-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention r thrombopoietin be used in the
                                                                                                                                                                                                                                                              Modified antibody; thrombopoietin; TPO; agonist; TPO receptor; platelet reduction-associated blood thrombocytopenia; cancer chemotherapy; leukaemia;
                                                                                                                            22-OCT-2001;
                                                                                                                                                           25-APR-2002
                                                                                                                                                                                          WO200233072-A1.
                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    used for improving neuromuscular function in a dystrophy. The products can also be used for cantibodies have a longer half-life than the nasequences AAX32387-X32413 represent DNA fragme CDR2, and CDR3 regions of variable heavy (VH) (VL) chains of antibodies Abl to Ab6.
                                                                                                                                                                                                                                                                                                                             Thrombopoietin (TPO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          thrombopoietin receptor agonist antibodies -
sting immunological or hematological disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10; Page 78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      They
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             etin receptor (TPO-R). The antibody (Ab) which binds to a the same way and for the same indications as thrombopoietin y can stimulate proliferation, differentiation or growth of ytes. They may also be able to stimulate megakaryocytes to latelet production. They can be used for treating cal or hematopoietic discreta.
                                                            2000JP-0321821.
2001WO-JP03288.
2001JP-0277314.
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 Tsunoda
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               unction in a patient, e.g. in muscular used for detection and diagnosis. The than the natural ligand for the TPO-R. DNA fragments encoding the CDR1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0 other;
                                                                                                                                                                                                                                                                                                                             associated
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signal
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hemotherapy for
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RESULT 3
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AC ABA045
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DT 115-FEE
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Best Loo
Matches
                                                     Antibodies for proliferation, activity to si
                                                                                                                                                                                      17-APR-2000;
20-OCT-2000;
20-OCT-2000;
12-MAR-2001;
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                                                                                                                                   Fukushima
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                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide
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                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABA04558;
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leukaemia
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                                                                                                                                                                                                                                                                                                                                                   .c; antiinflammatory; antianaemic; vasotropic; signal transmission; cancer; inflammation; hormonal; lymphoma; aplastic anaemia; skeletal malformation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            thrombopoietin agonist antibodies containing H and L chain V f monoclonal antibody, useful in preventives and/or remedies diseases, thrombocytopenia following cancer chemotherapy or
                            Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page
                                                     ignal transmission across
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2000JP-0321821.
2000JP-0321822.
2001WO-JP01912.
                                                  treatment of diseases associated with cell hormonal disorders and cytokines comprise rnal transmission across cell membranes -
                                                                                                                                                                                                                                                                                                                                                                                                                                    first
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                             157;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15; DB Pred. No. 55;
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                                                                                                                                 Ohtomo T;
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                                                                                                                                                                                                                                                                                                                                                                   disorder;
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invention

relates

to modified antibodies.

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antibodies

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RESULT 4
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Best Loo
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-OCT-2000;
17-APR-2001;
12-SEP-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hormonal disorders, leub
skeletal malformations.
the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene;
                                    The invention describes a modified antibody comprising at least 2 chain variable domains and 2 or more light chain variable domains antibody, and exhibits thrombopoietin (TPO) agonistic effect by cather the TPO receptor to crosslink. The antibodies are useful in preventand/or remedies for platelet reduction-associated blood diseases, thrombocytopenia following cancer chemotherapy or leukaemia. The antibody can act as a TPO signal transduction agonist by transductions signal into cells by crosslinking a TPO receptor to exert TPO agonist sequence encodes a thrombopoietin (TPO) agonist antibody associated by the control of the cont
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CHUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Modified antibody; thrombopoietin; TPO; agonist; TPO receptor; platelet reduction-associated blood thrombocytopenia; cancer chemotherapy; leukaemia;
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                                                                                                                                                                                                                                                                                  Degraded thrombopoietin agonist antibodies containing domains of monoclonal antibody, useful in preventives for blood diseases, thrombocytopenia following cancer leukaemia -
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P-PSDB; ABG35320.
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                                                                                                                                                                                                                                              Page 175-176;
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2001WO-JP03288.
2001JP-0277314.
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RESULT 5
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Best Loc
Matches
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20-OCT-2000;
20-OCT-2000;
12-MAR-2001;
                                               the antibodies care useful for twhich signal tra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; cyt antibody;
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P-PSDB;
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CDS
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                 hormonal
skeletal
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                                                                              of a monoclonal across the cell
                                                                                                                                                     proliferation activity to s
                                                                                                                                                                       Antibodies
                                                                                                                                                                                                                          Fukushima
                                                                                                                                                                                                                                                                                                                      17-APR-2001;
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)B; AAM47640.
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15; Cons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytostat
                sent invention relates to modified antibodies. The antibodies two or more H chain V domains and two or more L chain V domains noclonal antibody (MAb) which is capable of transmitting a signal the cell membrane by cross-linking a cell surface molecule, where ibodies can serve as signal transmission agonists. The antibodies ful for treatment and prevention of a broad range of disorders in ignal transmission is implicated, such as cancer, inflammation, l disorders, leukaemia, malignant lymphoma, aplastic anaemia, and l malformations. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                               CHUGAI
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ACCTACO
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2000JP-0321821.
2000JP-0321822.
2001WO-JP01912.
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                                                                                                                                                                                                                          Tsuchiya M,
                                                                                                                                                                                                                                               SEIYAKU KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ic; antiinflammatory; antianaemic; vasotropic; l transmission; cancer; inflammation; hormonal homa; aplastic anaemia; skeletal malformation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GGCATGCAC 15
                                                                                                                                                    treatment of diseases associated with cell hormonal disorders and cytokines comprise agonist nal transmission across cell membranes -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         irst entry)
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/*tag=
                                                                                                                                155-156; 173pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                     /partial
                                                                                                                                                                                                                                                                                                                                                                                  /product= "Human MPL #1"
/note= "No start or stop codon given"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 15; DB
100.0%; Pred: No. 58;
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                                                                                                                                                                                                                             Kikuchi Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
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Query Best 1

Match

Simila

city

100.0%; 100.0%;

Score Pred.

15; No.

DB 58;

24;

Length 351;

Sequence

351

ВP;

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73 T;

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ABA04563
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Best Local S
Matches 15
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17-APR-2001;
12-SEP-2001;
                                                                                                                                                                                                                                                              WPI; 200
P-PSDB;
                                                                                                                           signal
This se
                                                                                                                                                                                                                                                                                                                                                                                                                     Modified antibody; thrombopoietin; TPO; agonist; TPO receptor; platelet reduction-associated blood thrombocytopenia; cancer chemotherapy; leukaemia;
                                                                                                                                 The invention describes a modified antibody comprising at least 2 heavy chain variable domains and 2 or more light chain variable domains of an antibody, and exhibits thrombopoietin (TPO) agonistic effect by causing the TPO receptor to crosslink. The antibodies are useful in preventives and/or remedies for platelet reduction-associated blood diseases, thrombocytopenia following cancer chemotherapy or leukaemia. The antibody can act as a TPO signal transduction agonist by transducing a signal into cells by crosslinking a TPO receptor to exert TPO agonism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK71369;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK71369
                                                                                                                                                                                                                              Degraded thrombopoietin agonist antibodies containing domains of monoclonal antibody, useful in preventives for blood diseases, thrombocytopenia following cancer
                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA
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ABA04563
                                                                                                    Sequence
                                                                                                                   protein
                                                                                                                                                                                                                                                                                                                                                  22-OCT-2001;
                                                                                                                                                                                                        Example 7; Page 181-182; 213pp; Japanese
                                                                                                                                                                                                                        leukaemia
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)B; ABG35323.
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                                                                  Similarity 100.0%;
15; Conservative (
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                                       ACCTACGGCATGCAC
standard;
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2001WO-JP03288.
2001JP-0277314.
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DNA;
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Pred. No.
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No. 58;
matches
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                                                                                                                           agonist
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                                                                                    Length 432;
                                                                                                                                                                                                                                                                                                                                                                                                                       signal transduction;
                                                                                                                                                                                                                                                                                                                                                                                                                               disease;
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and/or remedic
chemotherapy
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RESULT 8
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11-MAY-1999
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Query Match
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Matches 15
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20-OCT-2000;
20-OCT-2000;
12-MAR-2001;
                                                                                                                                                                                                   The present invention relates to modified antibodies. The antibodies contain two or more H chain V domains and two or more L chain V domains of a monoclonal antibody (MAb) which is capable of transmitting a signal across the cell membrane by cross-linking a cell surface molecule, where the antibodies can serve as signal transmission agonists. The antibodies are useful for treatment and prevention of a broad range of disorders in which signal transmission is implicated, such as cancer, inflammation, hormonal disorders, leukaemia, malignant lymphoma, aplastic anaemia, and skeletal malformations. The present sequence was used to illustrate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; cytostatic; antiinflammatory; antianaemic; vasotropic; antibody; signal transmission; cancer; inflammation; hormonal disorder; leukaemia; lymphoma; aplastic anaemia; skeletal malformation; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fukushima N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-FEB-2002
                                                                                                                                                   Sequence 432
                                                                                                                                                                                                                                                                                                                                                                                                                                       activity to
                                                                                                                                                                                                                                                                                                                                                                                                                                                         proliferation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-OCT-2001.
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)B; AAM47642.
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15; Cor
                                    1 ACCTA
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ACCTA
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2000JP-0321821.
2000JP-0321822.
2001WO-JP01912.
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                                                                     100.0%; arity 100.0%; onservative 0
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                                                                                                                                                   P; 87
 CGGCATGCAC
                                    CGGCATGCAC 15
                                                                                                                                                                                          ion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEIYAKU KK.
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12..419
/*tag=_a
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/note= "No stop codon given"
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                                                                                                                                                   Α;
                                                                                                                                                   106 C;
   173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ID 63.
                                                                             ,;
                                                                        Score 15; DB Pred. No. 58; Mismatches
                                                                                                                                                   139 G;
                                                                                                                                                   100 T;
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                                                                                                                BG
                                                                                                            24;
                                                                                                                                                     0 other;
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                                                                             0;
                                                                                                               Length 432;
                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ohtomo T;
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                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                        This sequence represents the coding sequence of the anti-Mpl scFv antibody 12B5. The sequence encoding the chain is generated by a new method for preparing a multispecific Ab comprising a first polypeptide (PP) and at least 1 extra PP, where: (i) the first PP comprises a multimerisation domain (MD) forming an interface positioned to interact with an interface of a MD of the extra PP; and (ii) the first and extra PP seach have a binding domain, which comprises a heavy chain and a light C chain, where the variable light chains of the first and extra PP comprise a common sequence. The method comprises: (a) culturing a host cell comprising nucleic acid encoding the first PP and extra PP, and the variable light chain, such that the nucleic acid is expressed; and (b) recovering the multispecific Ab from the culture. The method prepares heteromultimeric PPs, such as bispecific Abs, bispecific immunoadhesins chaeromultimers and homomultimers. The method allows for the enhanced heteromultimers and homomultimers. The Abs can be used in immunoassays and for the in vitro or in vivo diagnosis of various diseases, such as
                                                                                                                                                                                                                                          Query Ma
Best Loo
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Variant; antibody;
diagnosis; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-JUN-1997;
02-MAY-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       selective prepara
heteromultimeric
useful for, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9850431-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anti-Mpl
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                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Arathoon R,
Modified antibody; thrombopoietin; TPO; agonist; TPO receptor; platelet reduction-associated blood thrombocytopenia; cancer chemotherapy; leukaemia; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example
                                                           DNA encoding
                                                                                   30-JUL-2002
                                                                                                         ABK71387
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timeric heavy chain and common light chain
r, e.g. in vivo diagnosis of cancer
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17-APR-2001;
12-SEP-2001;
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the TPO re
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Degraded thrombopoietin agonist antibodies containing domains of monoclonal antibody, useful in preventives for blood diseases, thrombocytopenia following cancer
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                                                                                                                                                                                      enase 12; cancer; HIV infection; enzyme;
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                                                                                                      AAA55186 to AAA55226 and AAY9091; and proteins isolated from the nc Cenarchaeum symbiosum. The nuclei the present invention are useful the present invention are used in the pre
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techniques
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Cenarchaeum
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physiology of
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P-PSDB; AAY90916
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ae; therapeutic; industrial; laboratory; ds.
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RESULT 13
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                                                                                                      AAA55186 to AAA55226 and AAY90913 to AAY90951 represent nucleic acids and proteins isolated from the non-thermophilic crenarchaeote Cenarchaeum symbiosum. The nucleic acids and proteins identified in the present invention are useful in characterising the physiology of these archae and can be used in therapeutic, industrial or laboratory techniques. AAA55227 to AAA55260 represent promoter sequences from Cenarchaeum symbiosum. AAA55261 to AAA55269 represent PCR primers and
                                                                              Sequence
                                                                                                                                                                                                                      crenarchaeote
physiology of
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ae; therapeutic; industrial; laboratory; ds.
13390
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Pred. No.
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                                      ore 14; DB 21;
red. No. 2.6e+02;
Mismatches 0;
                                                                                                 present invention.
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2.3e+02;
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RESULT 14
ABL71855,
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                                                                                                                                                                                                                                           quantitatively characterising the corresponding transcriptome, and in detecting RNA transcripts and splice variants of human or animal transcriptomes. The libraries may also be used as specialised minical transcriptomes. The libraries may also be used as specialised minical transcriptomes. The libraries may also be used as specialised minical transcriptome under a specific transcript of a sub-transcriptome under a graticular biological or pathological state, and so allowing the detection of tissue and pathology-specific genes such as those genes condition; to detect developmental specific genes; and to detect RNA transcripts and splice variants of a transcriptome of a patient suffering from a particular disorder. ABN27253 to ABN59589 represent oligonucleotide sequences from rats, humans and mice, which are used in the exemplification of the present invention.

Constructed the present invention.

The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at figure 1900 int/pub/published_pct_sequences.
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Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 several oligonucleotides, each capable of hese of messenger RNAs transcribed from a githe genome, which encodes one or more messenger the oligonucleotide libraries are useful for biological sample, in expression profiling quantitation.
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ABL71855
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02-MAY-2001;
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RESULT 15
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                                                                                                                                                                The present sequence describes a purified corn tassel-derived C polynucleotide sequence (cdps) comprising a nucleic acid sequence selected from those given in ABL70627 to ABL76833. The cdps sequences cencode corn tassel-derived polypeptides (CDPs). The cdps sequences (I) can be used for determining altered gene expression, to recover regulatory elements and to follow inheritance of desirable characteristics through hybrid breeding programs. (I) are also useful in the evaluation, and alteration of desired characteristics associated with growth and development, disease resistance, environmental adaptability, quality and yield, and as molecular markers for studying inheritance of multigene traits in a plant breeding program. (I) can be used to produce a tassel-specific profile of gene transcription, a transcript image, to clone regulatory elements for use in transformation vectors, to express a polypeptide, to identify, isolate or extend identical or related corn tassel nucleic acid sequences from DNA clibraries, in nucleic acid hybridisation or amplification technologies, as query sequences to determine homology of known sequences, as probe for use in Southern or Northern hybridisation, and to identify the presence of and/or to determine the degree of similarity between two
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93.3%;
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Pred. No. 4.6e+02;
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Search completed: June Job time: 88 secs
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                                                                                                                                                                              The present invention describes an isolated Rh(D) binding protein, preferably a human antibody, (I) having an amino acid sequence comprising one of the sequences (S) given in AAG93558 to AAG93669. (I) has immunostimulant activity, and can be used as an immune system stimulant. (I) can be used in diagnostic and therapeutic medicine. The antibodies are used in diagnostics that require human antibodies instead of animal antibodies, such as determine the Rh phenotype of human red blood cells. AAH68615 to AAH68726 represent the nucleotide sequence which encode AAG93558 to AAG93669. AAG93670 to AAG93697 represent anti-Rh(D) heavy chain CDR3 amino acid sequences which are given in the exemplification of the present invention.
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10-APR-1998;
27-JUN-1997;
                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                     New isolated protein, preferably a human anti-Rh(D) antibody for use diagnostics requiring a human instead of an animal antibody and in therapeutic medicine -
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P-PSDB; AAG93561.
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3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
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US-09-673-018-6
PCT-US95-09114-18
US-08-964-268-18
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EARLIER APPLICATION

EARLIER APPLICATION NUMBER: 60/020

EARLIER FILING DATE: 1996-10-11

NUMBER OF SEQ ID NOS: 224

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 78

PMGTH: 375
                                                               Patent No. 6255455

GENERAL INFORMATION:
APPLICANT: Siegel, Donald L.
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS !
TITLE OF INVENTION: SORTING METHOD FOR PRODUCE FILE REFERENCE: 09596-4202
CURRENT APPLICATION NUMBER: US/09/240,274
CURRENT FILING DATE: 1999-01-29
EARLIER APPLICATION NUMBER: 60/081,380
EARLIER APPLICATION NUMBER: 60/028,550
EARLIER FILING DATE: 1996-10-11
NUMBER OF SEQ ID NOS: 224
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APPLICANT: Siegel, Donald L.

TITLE OF INVENTION: Rh(D)-BINDING PROTEIN

TITLE OF INVENTION: SORTING METHOD FOR PR

FILE REFERENCE: 09596-42U2

CURRENT APPLICATION NUMBER: US/09/240,274

CURRENT FILING DATE: 1999-01-29

EARLIER APPLICATION NUMBER: 60/081,380

EARLIER FILING DATE: 1998-04-10
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GENERAL INFORMATION:
APPLICANT: Siegel,
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; FEATURE:
; NAME/KEY: CDS
; LOCATION: (36)..
US-09-388-743-5
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US-08-975-762-43
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US-09-3
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; OTHER INFORMATION US-09-240-274-93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 5
                                                                                                                                                                                                                                                                                   Sequence 43, Appli
Patent No. 6207169
GENERAL INFORMATI
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Best Local S
Matches 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/388,743
CURRENT FILING DATE: 1999-09-02
NUMBER OF SEQ ID NOS: 28
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: No. 6423886el Starch Synthase Polynucleotides TITLE OF INVENTION: Use in the Production of New Starches FILE REFERENCE: 1144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Singletary, George APPLICANT: Zhou, Lan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 23
                                                                                                       ZIP: 98104
COMPUTER READABL
MEDIUM TYPE:
                                                                                                                                                                                                TITLE OF INVENTION: (
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRES:
ADDRESSEE: SEED and
                                                                                                                                                                                                                                                 APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM:
ATTORNEY/AGENT NAME: Maki,
                                                          CURRENT APPLICATION DATA:
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uence 5, Applic
ent No. 6423886
                                                                                                                                                          STREET: 6300
CITY: Seattle
STATE: Washir
                      APPLICATION NUMBER: FILING DATE: 21-MA CLASSIFICATION: 42
                                                                      OPERATING SYS
SOFTWARE: Pa
                                                                                               COMPUTER:
                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                            868
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14; Conservative
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Lodes, Michael J.
Houghton, Raymond
VENTION: COMPOUNDS
                                                                  ADABLE FORM:

PE: Floppy disk

IBM PC compatible

SYSTEM: PC-DOS/MS-DOS

PatentIn Release #1.0, V
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illarity 93.3%;
Conservative
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SEED and BERRY LLP
Columbia Center,
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                                               US/08/975,762
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Pred. No. 93;
0; Mismatches
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Pred. No. 87;
0; Mismatches
                                                                                                                                                                                                                                     AND METHODS
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                                                                       Version
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David J.

INFORMATION:

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; STRANDEDNESS:
; TOPOLOGY: lin;
; MOLECULE TYPE:
US-08-975-762-43
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; ORGANISM: Ehrlichia
US-09-295-028-43
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US-09-295-028-43
                                                                                                                                                                                                           RESULT 8
US-09-106-582-43
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APPLICANT: Reed, Steven G.

APPLICANT: Lodes, Michael J.

APPLICANT: Houghton, Raymond L.

APPLICANT: McNeill, Patricia D.

TITLE OF INVENTION: COMPOUNDS AND ME
TITLE OF INVENTION: AND TREATMENT OF
FILE REFERENCE: 210121.439C4

CURRENT APPLICATION NUMBER: US/09/29

CURRENT FILING DATE: 1999-04-20

NUMBER OF SEQ ID NOS: 85
                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ :
SOFTWARE: Fast:
SEQ ID NO 43
LENGTH: 3735
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Best Local S
Matches 14
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                                                                                                                                                                      Patent No. 6306402
GENERAL INFORMATION:
                                                                                                                                                                                               Sequence
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                              APPLICANT:
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TELEPHONE: 206-622-4900
TELEFAX: 206-682-6031
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                               NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                       APPLICANT: Houghton, Raymond TITLE OF INVENTION: COMPOUNDS
                                  STREET:
CITY: Se
STATE: W
COUNTRY:
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                           ZIP:
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10. 6306402
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5. 6277381
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Lodes, Michael
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NUMBER: 21
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93.3%;
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OF EHRLI
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Mismatches
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RESULT 9
US-09-453-702B-57/c
; Sequence 57, Application US/09453702B
; Patent No. 6365723
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Matches 14
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INFORMATION FOR SEQ I
SEQUENCE CHARACTERI
LENGTH: 3735 bas
TYPE: nucleic ac
                                                               INFORMATION FOR
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-622-4900
TELEFAX: 206-682-6031
FORMATION FOR SEC. TO TELEPHONE SEC. TELEPHONE 
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CLASSIFICATION:
ATTORNEY/AGENT IN
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APPLICATION NUMBER: US/09/106 <0 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
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                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 60/110,955
FILING DATE: 04-DEC-1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 53701-2113
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch.
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 8.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: NO. NUMBER OF SEQUENCES: 26 CORRESPONDENCE ADDRESS: ADDRESSEE: Quarles
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                                     SEQUENCE
                                                                                                                                                            TELECOMMUNI
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NAME: Seay, Nicholas J.
REGISTRATION NUMBER: 27386
REFERENCE/DOCKET NUMBER: 960
ECOMMUNICATION INFORMATION:
TELEPHONE: (608) 251-5000
TELEFAX: (608) 251-9166
ION FOR SEQ ID NO: 57:
UENCE CHARACTERISTICS:
LENGTH: 87563
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Welch, Rod
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                                                                                                                                                                                                                                                                                                                                                                         CATION DATA:
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1 South Pinckney Street
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Pred. No. 9
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RESULT 11
PCT-US95-09114-18
Sequence 18, Application
GENERAL INFORMATION:
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US-09-673-018-6/c
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; SEQUENCE
US-09-453-702B-57
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: HASEGAWA, Junzo
APPLICANT: TAKAHASHI, Satomi
TITLE OF INVENTION: METHOD FOR PREPARING COI
FILE REFERENCE: 2000-1379A/LC/00177
CURRENT APPLICATION NUMBER: US/09/673,018
CURRENT FILING DATE: 2000-10-10
PRIOR APPLICATION NUMBER: JP 11/32657
PRIOR FILING DATE: 1999-02-10
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                   SOFTWALL SEQ ID NO 6
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Best Local
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                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                            TITLE
                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                      NUMBER OF
                                                                                                                                                       TITLE
                                                                                                                                                                  APPLICANT
                                                      STREET:
CITY: BC
STATE: M
COUNTRY:
                                            ZIP:
                                                                                                ADDRESSEE:
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NT: MATSUDA, Hideyuki
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Similarity 93.3%;
14; Conservative
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                                                                           Boston
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                                                                                                                     INVENTION:
INVENTION:
INVENTION:
F SEQUENCES:
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ULE TYPE: DNA (genomic)
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                                                                 Massachusetts
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IKENAKA, Yasuhiro
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compatible PC-DOS/MS-DOS
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92.9%;
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Mismatches
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1.1e+02;
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Binding
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US-08-990-823-15/c
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                                                                                                                                                                    SOFTWARE: Pat
SEQ ID NO 15
LENGTH: 265
TYPE: DNA
                                                  Best Loc
Matches
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Best Local
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                                                                                                     FEATURE:
NAME/KEY: Modified
OTHER INFORMATION:
-08-990-823-15
                                                                                                                                                                                                                                                                                                                                                                                           Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                    TITLE OF INVENTION: Mycobacterium Tuberculosis DNA Sequences
TITLE OF INVENTION: immunostimulatory Peptides
FILE REFERENCE: 49086
CURRENT APPLICATION NUMBER: US/08/990,823D
CURRENT FILING DATE: 1997-12-15
EARLIER APPLICATION NUMBER: US 96/10375
EARLIER FILING DATE: 1996-06-14
EARLIER APPLICATION NUMBER: 60/000,254
EARLIER FILING DATE: 1995-06-15
NUMBER OF SEQ ID NOS: 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEFAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                        ORGANISM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 08/305,903
FILING DATE: 13-Sep-94
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: DeConti, Giulio A., Jr.
REGISTRATION NUMBER: 31,503
REFERENCE/DOCKET NUMBER: BIZ-013CPPC
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/09114
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                                                                Local
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CGTAC
                        CCTACGGCATGCAC 15
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ilarity 92.9%;
Conservative
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92.9%;
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                                                Score 12.4; DB 4;
Pred. No. 3e+02;
Mismatches 1
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Sequence 182, Application US/09240274

Patent No. 6255455

GENERAL INFORMATION:
APPLICANT: Siegel, Donald L.
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACT
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
FILE REFERENCE: 09596-42U2

CURRENT APPLICATION NUMBER: US/09/240,274

CURRENT FILING DATE: 1999-01-29

EARLIER APPLICATION NUMBER: 60/081,380

EARLIER APPLICATION NUMBER: 60/028,550

EARLIER FILING DATE: 1996-10-11
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US-08-964-268-18
; Sequence 18, Ap
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APPLICANT: WL-
APPLICANT: RUBEN, J.
APPLICANT: SANCAR, AZIZ
APPLICANT: HSU, SHIAO-WEN D
APPLICANT: KAZANTSEV, ALEKSEY G
TITLE OF INVENTION: HUMAN BLUE-LIGHT PHOTOKE,
UMMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FO)
THIO NEW YORK AVENUE, NW, SUITE 60
THINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; MOLECULE TYPE: US-08-964-268-18
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Best Local S
Matches 13
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GENERAL INFORMATION:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/964,268
FILING DATE: HEREWITH
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/030,189
FILING DATE: 04-NOV-1996
ATTORNEY/AGENT INFORMATION:
NAME: STEFFE, ERIC K.
REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 1488.07500
TELEPHONE: 202-371-2540
TELEPAX: 202-371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 302 base pairs
TYPE: nucleic acid
                         NUMBER OF SEQ
            SOFTWARE:
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MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
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TOPOLOGY: li
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PatentIn
182
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Pred. No. 3
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RESULT 15

US-09-513-783A-45
; Sequence 45, Application US/09513783A
; Patent No. 6416959
; GENERAL INFORMATION:
; APPLICANT: Giuliano, Kenneth A.
; APPLICANT: Kapur, Ravi
; TITLE OF INVENTION: A System for Cell Based Screening
; FILE REFERENCE: 97-022-L1
; CURRENT APPLICATION NUMBER: US/09/513,783A
; CURRENT FILING DATE: 2000-02-25
; NUMBER OF SEQ ID NOS: 180
; COFTWARE: PatentIn Ver. 2.0
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3: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
4: /cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
5: /cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
6: /cgn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
7: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
9: /cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
10: /cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
11: /cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
12: /cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
13: /cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
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Sequence 1229, Ap

Sequence 73, Appl

Sequence 93, Appl

Sequence 612, Appli

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ALIGNMENTS

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RESULT 2
US-10-034-623-9
; Sequence 9, Application US/10034623
; Publication No. US20020198365A1
; GENERAL INFORMATION:
; APPLICANT: Swanson, Ronald V.
; APPLICANT: Feldman, Robert A.
; APPLICANT: Schleper, Christa
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Patent No. US20020160476A1
GENERAL INFORMATION:
APPLICANT: Swanson, Ronald V.
APPLICANT: Schleper, Christa
TITLE OF INVENTION: NUCLEIC ACIDS AND PROTEINS FROM CENARCHAEUM SYMBIOSUM
FILE REFERENCE: DCORP.002A
CURRENT APPLICATION NUMBER: US/10/027,806
CURRENT FILING DATE: 2001-12-21
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/408,020
PRIOR FILING DATE: EARLIER FILING DATE: 1999-09-29
NUMBER OF SEQ ID NOS: 123
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LENGTH: 15
TYPE: DNA
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NAME/KEY: CDS
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CURRENT APPLICATION NUMBER: US/10/U34, CURRENT FILING DATE: 2001-12-21

PRIOR APPLICATION NUMBER: 09/408,020

PRIOR FILING DATE: 1999-09-29

PRIOR APPLICATION NUMBER: 60/102,294

PRIOR FILING DATE: 1998-09-29

NUMBER OF SEQ ID NOS: 123

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; LOCATION: (1)...(1575)
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GENERAL INFORMATION:
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APPLICANT: Feldman, Robert A. APPLICANT: Schleper, Christa TITLE OF INVENTION: NUCLEIC ACIDS FILE REFERENCE: DCORP.002A CURRENT APPLICATION NUMBER: US/10/
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CURRENT FILING DATE: 2001-12-21
PRIOR APPLICATION NUMBER: EARLIER APPLICATION PRIOR FILING DATE: EARLIER FILING DATE: 1
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APPLICANT: Schleper, Christa
TITLE OF INVENTION: NUCLEIC ACIDS AND PROTEINS FROM CENARCHAEUM SYMBIOSUM
FILE REFERENCE: DCORP.002A
CURRENT APPLICATION NUMBER: US/10/034,623
CURRENT FILING DATE: 2001-12-21
PRIOR APPLICATION NUMBER: 09/408,020
PRIOR FILING DATE: 1999-09-29
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PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/408,020
PRIOR FILING DATE: EARLIER FILING DATE: 1999-09-29
NUMBER OF SEQ ID NOS: 123
SOFTWARE: FastSEQ for Windows Version 3.0
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NOS: 123
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US-10-027-801-2
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APPLICANT: Schleper, Christa
TITLE OF INVENTION: NUCLEIC ACIDS AND PROFILE REFERENCE: DCORP.002A
CURRENT APPLICATION NUMBER: US/10/027,801
CURRENT FILING DATE: 2001-12-21
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OTHER INFORMATION: Incyte ID No. US20010051335A1 700343977H1; NAME/KEY: unsure; LOCATION: 52, 62, 68, 83, 90, 145; OTHER INFORMATION: a, t, c, g, or other US-09-294-093B-1229
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LENGTH: 292
TYPE: DNA
ORGANISM: Zea may
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APPLICANT: Sherman, Bradley, K.
APPLICANT: Sherman, Bradley, K.
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN TASSEL
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN TASSEL
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN TASSEL
TITLE OF INVENTION: NUMBER: US/09/294,093B
CURRENT APPLICATION NUMBER: US/09/294,093B
CURRENT FILING DATE: 1999-04-16
PRIOR APPLICATION NUMBER: 60/082,567
                                                                                                                APPLICANT: Siegel, Donald L.
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
FILE REFERENCE: 09596-42U2
CURRENT APPLICATION NUMBER: US/09/848,798
CURRENT FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/240,274
PRIOR FILING DATE: EARLIER FILING DATE: 1999-01-29
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/028,550
PRIOR FILING DATE: EARLIER FILING DATE: 1996-10-11
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b. US20010051335A1
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Similarity 100.0%;
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                                                                                                                                                                                                                                                                                                            Application US/09848798
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PERL Program
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, Laura, Y.
                             sapiens
                                                                                                               E: EARLIER FILING IN NUMBER: EARLIER /
E: EARLIER FILING IN NOS: 224
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                                                                                                Ver.
anti-Rh(D) chain C04
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93.3%;
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Pred. No. 2e+02;
); Mismatches 1;
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75;
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US-09-848-798-73

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US-09-040 , ... Sequence 93, Application US/09040, ... Sequence 93, Application US/09040, ... Publication No. US20030040605A1 ; GENERAL INFORMATION:
APPLICANT: Siegel, Donald L.
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS A TITLE OF INVENTION: SORTING METHOD FOR PROD FILE REFERENCE: 09596-42U2
CURRENT APPLICATION NUMBER: US/09/848,798
CURRENT FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: EARLIER APPLICATIC PRIOR APPLICATION NUMBER: EARLIER APPLICATIC PRIOR APPLICATION NUMBER: EARLIER APPLICATIC PRIOR FILING DATE: 1999
NUMBER OF SEQ ID NOS: 224
NUMBER OF SEQ ID NOS: 224
                                                           ; OTHER INFORMATION: US-09-848-798-93
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US-09-848-798-93
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US-09-848-798-78
; Sequence 78, Application US/0984879
; Publication No. US20030040605A1
; GENERAL INFORMATION:
; APPLICANT: Siegel, Donald L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Дb
                    Best
    Matches
                              Query Match
                                                                                                                        NUMBER SOFTWARE: Partsey ID NO 93
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Best Local S
Matches 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: 09596-42U2
CURRENT APPLICATION NUMBER: US/09,
CURRENT FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: EARLIER
PRIOR FILING DATE: EARLIER FILING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORM 09-848-798-78
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TITLE OF INVENTION: Rh(D)-BINDING PHITITLE OF INVENTION: SORTING METHOD FILE REFERENCE: 09596-42U2
                                                                                  ORGANISM: Homo FEATURE:
                                                                                                                  TYPE:
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Similarity 93. 14; Conservative
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In Ver.
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93.3%;
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Score 13.4; In Pred. No. 2e+00; Mismatches
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DATE: 1999-01-29
APPLICATION NUMBER:
DATE: 1996-10-11
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DATE: 1999-01
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Mismatches
   13.4;
2e+02;
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No. 2e+02;
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APPLICANT: Halk, Edward

APPLICANT: Gerritsen, Arnout F.

TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO

TITLE OF INVENTION: EPIDERMAL GROWTH FACTOR RECEPTOR (EGFR)

FILE REFERENCE: GMI-020

CURRENT APPLICATION NUMBER: US/10/172,317

CURRENT FILING DATE: 2002-06-13

PRIOR APPLICATION NUMBER: US 60/298,172

PRIOR FILING DATE: 2001-06-13

NUMBER OF SEQ ID NOS: 4
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US-10-172-317-
SEQ
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Matches
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                    CURRENT APPLICATION NUMBER: US/09/822,846
CURRENT FILING DATE: 2001-03-29
PRIOR APPLICATION NUMBER: 60/195,605
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 629
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APPLICANT:
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TYPE: DNA
ORGANISM: Homo
-10-172-317-1
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             SOFTWARE:
                                                                                                          TITLE OF
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                                                                                          REFERENCE: GIN
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             Patent
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4; Conservative
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                                                                                                                                                                                   Agostino, Michael J.
Steininger II, Robert
Bowman, Michael R.
Spaulding, Vikki
Wong, Gordon G.
Clark, Hilary
Fechtel, Kim
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Resnick,
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                                                                                         Resnick, Richard J.
Gulukota, Kamalakar
Graham, James R.
Genetics Institute, Inc.
Genetics POLYNUCLEOTIDES ENCODING NOVEL SECRETED PROTEINS
ICE: GIN 6400
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Merberg
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berg, Maurice
arv, Maurice
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S20030027139A1
ON:
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TGGCATGCAC 105
             NOS: 6
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) for Windows Version 4.0
                                                                                                                                                                                                                                                                                                          Cheryl
                                                                                                                                                                       Steven H
                                                                                                                                                                                                                                                                                                                                               John M.
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michael R.
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93.3%;
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Pred. No. 2
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2e+02;
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Sequence 6, Application US/09816760

Patent No. US20020052032A1

GENERAL INFORMATION:
APPLICANT: Meyers, Rachel
APPLICANT: Cook, William James
APPLICANT: Williamson, Mark
APPLICANT: Rudolph-Owen, Laura A.
TITLE OF INVENTION: 32142, 21481, 25964, 21686, NOV
TITLE OF INVENTION: MOLECULES AND USES THEREFOR
FILE REFERENCE: MNI-134CP
CURRENT APPLICATION NUMBER: US/09/816,760
CURRENT FILING DATE: 2001-03-23
PRIOR APPLICATION NUMBER: 09/634,955
PRIOR APPLICATION NUMBER: 60/192,002
PRIOR FILING DATE: 2000-08-08
PRIOR FILING DATE: 2000-03-24
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US-09-838-561-6
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US-09-816-760-6
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; NAME/KEY: CDS
; LOCATION: (1)..(933)
US-09-838-561-6
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TITLE OF INVENTION: 32142, 21481, 25964, 21686, NOV

TITLE OF INVENTION: MOLECULES AND USES THEREFOR

FILE REFERENCE: MNI-134CP2

CURRENT APPLICATION NUMBER: US/09/838,561

CURRENT FILING DATE: 2001-04-18

PRIOR APPLICATION NUMBER: 09/816,760

PRIOR FILING DATE: 2001-03-23

PRIOR APPLICATION NUMBER: 09/634,955

PRIOR APPLICATION NUMBER: 09/634,955

PRIOR APPLICATION NUMBER: 60/192,002

PRIOR FILING DATE: 2000-08-08

PRIOR FILING DATE: 2000-03-24

NUMBER OF SEQ ID NOS: 16

SOFTWARE: Patentin Ver. 2.0
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Best Local S
Matches 14
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LENGTH: 933
TYPE: DNA
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APPLICANT: Meyers, Rachel

APPLICANT: Cook, William James

APPLICANT: Williamson, Mark

APPLICANT: Rudolph-Owen, Laura
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PRIOR FILING DATE: NUMBER OF SEQ ID N
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93.38;
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Patent No. US20020076/UDAL

GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: 31 Human Secreted Proteins
FILE REFERENCE: PZ033P1
CURRENT APPLICATION NUMBER: US/09/820,893
CURRENT FILING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: 09/531,119
PRIOR FILING DATE: 2000-03-20
PRIOR FILING DATE: 1900-03-20
PRIOR FILING DATE: 1998-10-02
NUMBER OF SEQ ID NOS: 140
SCOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; FEATURE:
; NAME/KEY: CDS
; LOCATION: (1)..(
US-09-816-760-6
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US-09-820-893-31
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Search completed:
Job time : 65.2 s
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; ORGANISM: Homo
US-09-820-893-31
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SOFTWARE: PatentII
SEQ ID NO 31
LENGTH: 1157
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TYPE: DNA
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V385001 AV38500	V38500	10	\sim	ω ·	14	20
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TITLE
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COMMENT
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VERSION
KEYWORDS
SOURCE
ORGANISM
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R12983/c
LOCUS
DEFINITION
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AUTHORS
                 1 (bases 1 to 354)
Hillier, L., Clark, N., Dubuque, T., Elliston, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., R., Williamson, A., Wohldmann, P. and Wilson, R., Williamson, A., Wohldmann, P. and Wilson The WashU-Merck EST Project
Unpublished (1995)
Contact: Wilson RK
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Lou
Tel: 314 286 1800
Fax: 314 286 1810
Email: est@watson.wustl.edu
Insert Size: 2106
High quality sequence stops: 175 Source: IN
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IMAGE: 2
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l: est@watson.wustl.edu
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quality sequence stops: 175 Source: IMAGE Consortium, LLNL
clone is available royalty-free through LLNL; contact the
E Consortium (info@image.llnl.gov) for further information.
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Proc. Natl. Acad. Sci. U.S.A. 97 (7),
20202663
Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 anda
                                                 Tel: +55-11-2704922

Fax: +55-11-2707001

Email: asimpson@ludwig.org.br

This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL (http://www.ludwig.org.br/scripts/gethtml2.pl?t1=QV3&t2=QV3-BN0046 060700-256-d10&t3=2000-07-06&t4=1)

Seq primer: puc 18 forward
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1 (bases 1 to 549)

Dias Neto, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R., Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F., Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H., Brunstein, A., deOliveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and
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Seq primer: M13
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BG539077/c
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Eukaryota;
Mammalia; E
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NIH-MGC ht
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Unpublished (1999)
Contact: Robert St
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BG539077
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602568449F1 NIH_MGC_77 Homo sapiens
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15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                        quality sequence stop: 534.
Location/Qualifiers
                                                                                    /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4692781"
/clone="IMAGE:4692781"
/clone=lib="NIH_MGC_77"
/lab_host="DH10B (T1 phage-resistant)"
/note="Organ: lung; Vector: pDNR-LIB (Clontech); Site_1:
Sfil (ggccgcctcggcc); Site_2: Sfil (ggccattatggcc); 5' and
3' adaptors were used in cloning as follows: 5' adaptor
sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor sequence:
5'-ATTCTAGAGGCCGAGGCGGCCGACATG-dT(30)BN-3' (where B = A,
C, or G and N = A, C, G, or T). Average insert size 1.9
kb (range 0.5-4.0 kb). 12/15 colonies contained inserts
by PCR. This library was enriched for full-length clones
and was constructed by Clontech Laboratories (Palo Alto,
CA). Note: this is a NIH_MGC Library."

46 a 131 c 178 g 208 t
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/db_xref="taxon:9606"
/clone_lib="BN0046"
/dev_stage="Adult"
/note="Organ: breast_normal; Vector: pucl8; Site_l: SmaI;
Site_2: SmaI; A mini-library was made by cloning products
derived from ORESTES PCR (U.S. Letters Patent application
No. 196,716 - Ludwig Institute for Cancer Research)
profiles into the pUC 18 vector. Reverse transcription of
tissue mRNA and cDNA amplification were performed under
low stringency conditions."
13 a 133 c 156 g 147 t
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Institutes of Health, Mammalian Gene Collection (MGC)
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Eutheria;
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                      100.0%;
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   Score 15; DB Pred. No. 1.6); Mismatches
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Pred. No.
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. 1.6e+03;
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hes 0;
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                                       Length 663;
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  (where B = A,
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Conservative

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Gaps

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29

GGCATGCAC

15

linear

EST 20-OCT-2000

Euteleostomi;

Homo

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BM716521/c
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Genome Res.
97044477
Contact: Soa
Program for
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Mammalia; E
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BM716521.1
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Seq prime
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UI-E-EJO-ahi-i-11-0-UI.r2 UI-E-EJO Homo UI-E-EJO-ahi-i-11-0-UI 5', mRNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: msoares@blue.weeg.uiowa.edu
Tissue Procurement: Dr. Gregg Hageman
cDNA Library preparation: Dr. M. Bento Soares, Univeristy of
cDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of
DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa
Clone Distribution: Researchers may obtain clones from Resear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 706)
Bonaldo, M.F., Lennon, G. and Soares,
Normalization and subtraction: two
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                  15;
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451 Eckstein Medical Research Building
Tel: 319 335 8250
Fax: 319 335 9565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human.
                                 Similarity
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                                                                                                   183
                  Conservative
                                                                                                        /dev_stage="fetal and adult"
/lab_host="DH10B (Life Technologies) (T1 phage resistant)"
/note="Organ: eye; Vector: pT7T3-Pac (Pharmacia) with a
modified polylinker; Site_1: Ecor I; Site_2: Not I;
UI-E-EJO is a subtracted cDNA library constructed
according to Bonaldo, Lennon and Soares, Genome Research,
6:791-806, 1996. First strand cDNA synthesis was primed
with an oligo-dT primer containing a Not I site. Double
stranded cDNA was ligated to an Ecor I adaptor, digested
with Not I, and cloned directionally into pT7T3-Pac
vector. The oligonucleotide used to prime the synthesis of
first-strand cDNA contains a library tag sequence that is
located between the Not I site and the (dT)18 tail. The
sequence tags for this library are: fetal eyes, AGAATCAAGA; eye anterior segment, AATGCCGCAT;
optic nerve, CCATTAAGTG; retina, CCGCG; Retina Foveal and
Macular, GTCC; RPE and Choroid, ACCTA. This library was
created for the program, Gene Discovery in the Visual
System, supported by National Eye Institute (NEI)."

130 C 228 t 1 others
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                                                                                                   Ø
                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="UI-E-EJO-ahi-i-11-0-UI"
/clone_lib="UI-E-EJO"
/tissue_type="fetal eyes, lens, eye anterior se optic nerve, retina, Retina Foveal and Macular, Choroid"
                                                                                 Macular, GTCC; KPL and Control of the program, Gene created for the program, Gene System, supported by National 130 c 164 g 228 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (www.resgen.com).
r: M13 Reverse.
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Eutheria; Primates; Catarrhini;
1 to 706)
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1.6e+03;
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b; Hominidae;
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                                              Length 706;
                                                                                                   1 others
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Matches 15
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                                                               EFERENCE
AUTHORS
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Eukaryota; Me
Mammalia; Eut
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mRNA sequence.
BE882505
BE882505.1 GI
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NIH-MGC ht
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601507246
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BQ719668
BQ719668
                                                                                             Homo sapi
Eukaryota
Mammalia
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AGENCOURT
 Contact: Email: c
                        Nim-MGC http://mgc.nci.nih.gov/
National Institutes of Health,
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov Plate: LLAM9721 row: l column: 12
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                                                                                                                                                                  EST.
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                                                                                                                                                  human.
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                                                                                                                                                                                                                                                                                                                                                                ACCTAC
                                                                                (base
                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
milarity 100.0%;
Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:3908891"
/clone_lib="NIH_MGC_71"
/tissue_type="leiomyosarcoma"
/lab_host="DH10B (phage-resistant)"
/note="Organ: uterus; Vector: pCMV-SPORT6; Site_1:
Site_2: Sall; Cloned unidirectionally. Primer: Ol.
Average insert size 2.1 kb.
0 a 228 c 255 g 270 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                t: Robert Strausberg, Ph.D. cgapbs-r@mail.nih.gov
Procurement: ATCC
: Robert Strausberg, cgapbs-r@mail.nih.go
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6F1 NIH_MGC_71 Homo sapiens cDNA clone IMAGE:3908891 5'
                                                                             a; Metazoa; Chordata; Craniata; Vertebrata; ; Eutheria; Primates; Catarrhini; Hominidae s 1 to 961)
                                                                                                                                iens
                                                                                                                                                                                                                 93699 5′,
                                                                                                                                                                                                                                                                                                                                                                GGCATGCAC 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                http://mgc.nci.nih.gov/.
Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ity sequence stop: 608.
Location/Qualifiers
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Eutheria;
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Primates;
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Pred. No. 1.8e+03;
Mismatches 0;
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Catarrhini;
                                                 Mammalian Gene Collection (MGC)
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i; Hominidae;
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Length 923;

Oligo dT. 1: NotI;

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16-JUL-2002 cDNA clone

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TITLE
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                                                                                                                                                                     Williams, B.A., Tsang, A. a Sequence analysis of 68 c Aspergillus niger cDNA li Unpublished (1995) Contact: Williams BA and Dept of Biology Concordia University
                                                                                                                                                                                                                                                                                                                            5' end, mRNA sequen
T82729
T82729.1 GI:711017
EST.
Aspergillus niger.
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                                                                                    (nov01t)
Seq prime
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Nig011
5' end,
                                                                                                                         1455 de Maisonneuve Blvd
Tel: (514) 848 3412
Fax: (514) 848 2881
                                                                                                                                                                                                                                                                      Aspergillus niger
Eukaryota; Fungi; Ascomycota;
Eurotiales; Trichocomaceae; mi
1 (bases 1 to 268)
                                                                                                                                                                                                                                                      Williams, B.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              . Similarity 15; Conserv
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                                                                                     primer:
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quality sequence stop:
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="Vector: pCMV-SPORT6 (Life Technologies); Site_1:
NotI; Site_2: SalI; cDNA made by oligo-dT priming.
Directionally cloned using the following adaptors:
5'-TCGACCCACGCGTCCG-3' and
5'-GACTAGTTCTAGATCGCGAGCGGCCCCT(15)-3'. Size selected >
1 kb for average insert length 1.9 kb. This is a primary
library, non-amplified. Library constructed by Life
Technologies and donated by J. Lupski, M.D./Ph.D. (Baylor College of Medicine); available through Life
Technologies."
37 a 274 c 248 a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                      Aspergillus niger, mRNA sequence.
                                                                                                                 storms@vax2.concordia.ca
                                                                  r: T7 Pro Primer.
Location/Qualifiers
/organism="Aspergillus r/db_xref="taxon:5061"
/clone_lib="Aspergillus
/lab_host="E. coli"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:6193699"
/clone_lib="Lupski_sympathetic_trunk"
/sex="male"
                                                                                                                                                                                                                                                                                                                                                           GI:711017
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age="adult, 16 yr"
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                                                                                                                                                                                                                             and Storms, R.K clones selected library
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                                                                                                                                                        West,
                                                                                                                                                                                                                                                                                                                                                                                                                 268
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): 583.
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mitosporic Trichocomaceae;
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ES2
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                                                                                                                                                                                                                                            selected
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             niger,
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t
              pYES2
                                                                                                                                                         Quebec,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 961;
                                                                                                                                                                                                                                           random
                                                                                                                                                                                                                                                                                               Eurotiomycetes;
              (XhoI-EcoRI)"
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AUTHORS
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SOURCE
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AI764493,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  University of
451 Eckstein M
Tel: 319 335 8
Fax: 319 335 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 (bases Bonaldo, M.)
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9704447
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AI76449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: msoares@blue.weeg.uiowa.edu
Oligo-dT track not found, Not I site shown in beginning of sequence
is likely internal to the message. cDNA Library Preparation: M.B.
Soares Lab Clone distribution: clones will be available through
Research Genetics (www.resgen.com) The following repetitive
elements were found in this cDNA sequence: 95-157, >(CGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel:
Fax:
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Normalization and subtraction: two approaches to facilitate gene
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Mammalia;
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.4; Conservative
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R-Y0-abk-a-10-0-UI 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          primer:
/dev_stage="adult"
/lab_host="DH10B (Life Technologies)"
/note="Vector: pT7T3D-Pac (Pharmacia) with a modified
/note="Vector: pT7T3D-Pac (Pharmacia) with a 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          norvegicus
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                                                                                                                                                                                                                                                                                                                                                                                             /organism="Rattus norvegicus"
/strain="Sprague-Dawley"
/db_xref="taxon:10116"
/clone="UI-R-Y0-abk-a-10-0-UI"
/clone_lib="UI-R-Y0"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       I to
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Pred. No. 3.6e+03;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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Query Match
Best Local
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                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammmalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus.

CE 1 (bases 1 to 369)

RS Caetano, A.R., Johnson, R.K. and Pomp, D.

Generation and sequence characterization of a normalized cDNA library from swine ovarian follicles

Unpublished (2001)

Contact: Pomp, D

Department of Animal Science
University of Nebraska, Lincoln

Lincoln, NE 68583-0908, USA

Tel: 402 472 6362

Email: dpomp@unl.edu

The sequence contained an oligo-dT track that was present in the oligonucleotide that was used to prime the synthesis of first strand cDNA and therefore this may represent a bonafide poly A tail. The sequence tag present in the cDNA between the NotI site and the oligo-dT track served to verify it as a clone from the normalized porcine ovarian follicles library The following repetitive elements were found in this cDNA sequence: 258-349, sqi|2087|emb|X64127.1|SSPRE S.scrofa DNA for SINE sequence SSPRE Seq primer: M13 -29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BI183951
BI183951.1
EST.
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Sus
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UNL-P-FN-cf-e-03-0-UNL.s1 UNL-P-FN Sus so
UNL-P-FN-cf-e-03-0-UNL 3', mRNA sequence
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TAG_TISSUE=Eye
TAG_SEQ=CATTG"
/db_xref="taxon:9823"
/clone="UNL-P-FN-cf-e-03-0-UNL"
/clone_lib="UNL-P-FN"
/clone_lib="UNL-P-FN"
/dev_stage="ADULT"
/lab_host="DH10B (Life Technologi/note="Vector: pT7T3D-Pac (Pharmapolylinker; Site_1: Not I; Site_2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amplified cDNA inserts from previous library clones in which 3' ESTs had been derived were used as a driver hybridization with the normalized whole-eye library if the form of single-stranded circles. The remaining single-stranded circles (subtracted library) was purify hydroxyapatite column chromatography, converted to double-stranded circles and electroporated into DH10B double-stranded circles and electroporated the UI-R-Y0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (Bonaldo,
1996)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bacteria (Life Technologies) to generate the UI-R-Y0 library. This procedure has been previously described (Bonaldo, Lennon and Soares, Genome Research 6: 791-806,
                                                                                                                                              /organism="Sus scrofa"
/strain="University of Nebraska,
Lines"
                                                                                                                                                                                                                                ocation/Qualifiers
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3.9e+03;
0;
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Pac (Pharmacia) with
t I; Site_2: Eco RI;
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he UNL-P-FN
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REFERENCE AUTHORS TITLE

COMMENT

JOURNAL

FEATURES

KEYWORDS SOURCE

ORGANISM

ACCESSION VERSION

RESULT 9 BI183951

DEFINITION

LOCUS

В

BASE COUNT ORIGIN

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Mammalia; E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14;
                                                                                                                                                                                                                                                                                                                                                                             Email: msoares@blue.weeg.uiowa.edu
Tissue Procurement: Dr. Gregg Hageman
cDNA Library preparation: Dr. M. Bento Soares, Univeristy
cDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy
DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iow
Clone Distribution: Researchers may obtain clones from Re
Genetics (www.resgen.com).
Seq primer: M13 Reverse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BM728351 486 bp mRNA linear EST 01 UI-E-E01-aiv-i-15-0-UI.rl UI-E-E01 Homo sapiens cDNA clone UI-E-E01-aiv-i-15-0-UI 5', mRNA sequence.
BM728351 ...
BM728351.1 GI:19049684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    University of Iowa
451 Eckstein Medical Research Building Iowa City,
Tel: 319 335 8250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: 319 335 8250
Fax: 319 335 9565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9704447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genome
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arity 100.0%;
onservative C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       library is a normalized library representing porcine ovarian follicles, ranging between 2.0 to 10.0 mm in diameter, collected during 7 days of the follicular phase of the pig estrous cycle. This library was derived from the library UNL-P-F2. The tag is a string of 5-6 nucleotides present between the Not I site and the oligo-dT track. The library was constructed as described by Bonaldo, Lennon and Soares, Genome Research 6: 791-80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TAG_LIB=UNL-P-FN
TAG_TISSUE=porcine ovarian follicles
TAG_SEQ=CACACT"
AA A BA B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Soares, MB
or Rat Gene Discovery and Mapping
following tissue(s): fetal eye. The library was constructed according to Bonaldo, Lennon and Soares, Genome Research, 6:791-806, 1996. First strand cDNA synthesis was primed with an oligo-dT primer containing a Not I site. Double stranded cDNA was ligated to an EcoR I
                                                                                             /organism="Homo sapiens"
/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="UI-E-E01-aiv-i-15-0-UI"
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/clone_lib="UI-E-E01"
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/clone_lib="UI-E-E01"
/clone_lib="UI-E-E01"
/tissue_type="fetal eye"
/dev_stage="fetal eye"
/dev_stage="fetal"
/lab_host="DH10B (Life Technologies) (T1 phage resistant)"
/note="Organ: eye; Vector: pT7T3-Pac (Pharmacia) with a modified polylinker; Site_1: EcoR I; Site_2: Not I;
UI-E-E01 is a normalized cDNA library containing the
                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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Pred. No.
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4.1e+03;
                                                                                            The library was
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JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                Email: est@watson.wustl.edu
Libraries were constructed by Dr. Stavros Bashiardes as part of the physicomitrella EST program (PEP) at the Univ. of Leeds (UK) and Washington Univ. in St. Louis (USA) DNA sequencing by: Washington University Genome Sequencing Center For information on obtaining clone please contact: Celia Knight (c.d.knight@leeds.ac.uk)
High quality sequence stop: 421.

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BQ040153
BQ040153.1
EST.
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Bryophyta;
Bryopsida; Funariidae; Funariales; Funariaceae; Physicomitrella.
Bryopsida; Funariidae; Funariales; Funariaceae; Physicomitrella.
(bases 1 to 531)
Quatrano,R., Bashiardes,S., Cove,D., Cuming,A., Knight,C., Clifton,S., Marra,M., Hillier,L., Pape,D., Martin,J., Wylie,T., Underwood,K., Theising,B., Allen,M., Bowers,Y., Person,B., Swaller,T.,
Steptoe,M., Gibbons,M., Harvey,N., Ritter,E., Jackson,Y., McCann,R., Waterston,R. and Wilson,R.
Leeds/Wash U Moss EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1999)
Other_ESTs: gdl2b08.xl
Contact: Ralph Quatrano
Leeds/Wash U Moss EST Project
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. I
Tel: 314 286 1800
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531 bp mRNA linear EST 17-APR-200: gd12b08.y1 Moss EST library PPS Physcomitrella patens cDNA clone PEP_SOURCE_ID:PPS40315 5' similar to TR:Q41050 Q41050 CORE PROTEIN
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the pBluescript sequence as well as the lambda DNA and CDNA is cloned in the EcoRI and XhoI sites in the pBluescript sequence. The vector was then packaged using Gold gigapackaging extracts, propagated in XL-IBlue MRF
                                                                       /note="Vector: pBluescript SK-; Site_1: XhoI; Site_2: ECORI; Library constructed by Stavros Bashiardes and re-arrayed by A. Cuming & Honglin Rong. Construction of the cDNA library was carried out using Statagene's 'Unizate CDNA directionally in Unizap XR vector arms. The vector is designed containing the pBluescript sequence as well as the lambda DNA and
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                                                                                                                                                                                                                                                                   /organism="Physcomitrella patens"
/db_xref="taxon:3218"
/clone="PEP_SOURCE_ID:PPS40315"
/clone_lib="Moss EST library PPS"
/dev_stage="protonemata, 7day old
/lab_host="E.coli DH10b"
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Pred. No. 4.5e+03;
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Sheared
Sheared
AQ657745
AQ657745
GSS.
                                                                                                                                                                                                                                                                                                                                                                                             Email: nelsayed@tigr.org
Clones are derived from the Trypanosoma brucei GUTat 10.1 sheared
DNA library constructed at TIGR. Clones will be available for
distribution through ATCC. Sheared DNA end sequences search page:
http://www.tigr.org/tdb/mdb/tbdb/.
Seq primer: M13-Forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fax:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Determination of clone end sequences from Trypanosoma brucei GUTat 10.1 sheared DNA library
Unpublished (1999)
Other_GSSs: Sheared DNA-27J6.TR
Contact: Najib M. El-Sayed
Department of Eukaryotic Genomics
The Institute for Genomic Research
9712 Medical Center Dr., Rockville, MD 20850, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tel:
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/Clone_lib="Sheared DNA"
/clone_lib="Sheared DNA"
/note="Vector: pUC18; Site_1: SmaI; Constructed at The
Institute for Genomic Research (TIGR), Rockville, MD.
Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically
sheared to give a tight size distribution (approx 2 kb).
The v + i method used for the library construction is
described in detail in Smith, H.O. and Venter, J.C.
(Making small insert libraries for whole genome shotgun
sequencing projects. In Genome Sequencing: A Practical
Approach, eds. M. Vaudin and B. Barell, Oxford University
Press, 1999)."
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838 0208
                                                                                                                                                                                                                                                  /organism="Trypanosoma brucei"/strain="TREU927/4 GUTat 10.1"/db_xref="taxon:5691"/clone="Sheared DNA-27J6"
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Leech,V.,
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RESULT 13
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L (bases 1 to 591)
El-Sayed, N., Zhao, S., Zhao
Gerrard, C., Leech, V., de of Fraser, C. and Adams, M.
Determination of clone end
10.1 sheared recommend
                                                                                                                                                                                                  160
                                                                                                                                                                                                                                                                                                                                                        Kyoto University
Sakyo-ku, Kyoto, Kyoto 606-850
Tel: 81-75-753-4095
Fax: 81-75-705-1113
Email: kwmakabe@ascidian.zool.
Location/Qualifiers
e 1. .569
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                                                                                               AV386289 Ha
CDNA clone
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Trypanosoma.
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Unpublished (1999)
Contact: Kazuhiro W. Makabe
Department of Zoology, Grad
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Urochord. Stolidobranchia; Pyuridae; Halocynthia 1 (bases 1 to 569)
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Similarity 100.0%;
14; Conservative
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Halocynthia roretzi
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                                                                                                                                                                                                                                                                                      /organism="Halocynthia rc/db_xref="taxon:7729"
/clone="009B23_3"
/clone_lib="Halocynthia rc/dev_stage="Fertilized ec/note="Organ: embryo"
a 126 c 124 g 162
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Contact:
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Clones are derived from the Trypanosoma brucei GUTat 10.1 sheared
Clones are derived from the Trypanosoma brucei GUTat 10.1 sheared
DNA library constructed at TIGR. Clones will be available for
distribution through ATCC. Sheared DNA end sequences search page:
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The Institute for Genomic Research
9712 Medical Center Dr., Rockville,
Tel: 301 838 0200
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http://www.tigr.org/tdb/mdb/tbdb/.
Seq primer: M13-Reverse
Class: shotgun.
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                                                                                             Department of Eukaryotic Genomics
The Institute for Genomic Research
9712 Medical Center Dr., Rockville
Tel: 301 838 0200
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Clones are derived from the mouse BAC library RPCI-23. For BAC library availability, please contact Pieter de Jong (pieter@dejong.med.buffalo.edu). Clones may be purchased from
                                                                               Fax:
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                                                                                                                                                                                                                                                                                                                  Mammalia
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                                                                                                                                                                                                                                                                                                                                                                         house mouse
                                                        Email: szhao@tigr.org
                                                                                                                                                                             Contact:
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Levins,M., Mcgann,S., Tsegaye,G., Geer,K., Krol,M., de Jong,P.
7raser,C.M.
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.larity 100.0%;
Conservative
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/note="Vector: pUCl8; Site_l: SmaI; Constructed at The
Institute for Genomic Research (TIGR), Rockville, MD.
Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically
sheared to give a tight size distribution (approx 2 kb).
The v + i method used for the library construction is
described in detail in Smith, H.O. and Venter, J.C.
(Making small insert libraries for whole genome shotgun
sequencing projects. In Genome Sequencing: A Practical
Approach, eds. M. Vaudin and B. Barell, Oxford University
Press, 1999)."
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itute for Genomic Research
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/clone="sheared DNA-53M15"
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/strain="TREU927/4 GUTat 10.1"
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Center Dr., Rockville,
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                                                          BACPAC Resources (http://bacpac.med.buffalo.edu/orderingframe.htm) or from Resea ch Genetics (info@resgen.com). BAC end page: http://www.tigr.org/tdb/bac_ends/mouse/bac_end_intro.html Plate: 21 row: H column: 20 Seq primer: SP6 Class: BAC ends.

Location/Qualifiers
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                                                                                                                                                                                                                       /organism="Mus musculus"
/organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="RPCI-23-21H20"
/clone="RPCI-23-21H20"
/clone="Pemale"
/lab_host="DH10B"
/note="Organ: Kidney/Brain; Vector: pBACe3.6; Site_1:
ECORI; Site_2: ECORI; Female C57BL/6J mouse kidney and/or
brain genomic DNA was isolated and partially digested
with a combination of EcoRI and EcoRI Methylase. Size
selected DNA was cloned into the pBACe3.6 vector at the
ECORI sites. The ligation products were transformed into
DH10B electrocompetent cells (BRL Life Technologies). "
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QY Search completed: June Job time: 754.2 secs Дb Query Match 93.3%; Best Local Similarity 100.0%; Matches 14; Conservative

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Unknown. M Unknown. Unclassified. 1 (bases 1 to 15) Adams, C.W., Carter, P.J., Fen Agonist antibodies Patent: US 6342220-A 36 29-J	AR183492 N Sequence 36 from patent US 6 AR183492 AR183492.1 GI:20227461	ALIG	3.3 64767 9 U69730 3.3 71320 2 AC09541	78 3 AC00433 31 2 AC11847	3.3 60328 9 AL50052	3.3 38109 1 MLCB5	3.3 23819 2 AC01817 3.3 36138 1 MLU1518	3.3 9908 1 AE01330	3.3 2898 1 LIU9505	3.3 2570 8 AF42217	3.3 652 11 PM7H5B	4 93.3 415 8 AY03811	93.3 258 9 HSAJ951 93.3 373 8 AY03811	5 100:0 22/010 2 ACT1349 4 93.3 127 9 HUMRB	5 100.0 221372 2 AC12447	5 100.0 205202 2 AC10879 5 100.0 207473 2 AC11763	5 100.0 182991 2 AC10165	5 100.0 180613 2 AC11506	5 100.0 170717 2 AC12859 5 100.0 175518 9 AC01884	5 100.0 165733 2 AC09914	5 100.0 154084 9 AC02231	5 100.0 145520 2 AC01603	5 100.0 127400 2 AC13112	5 100.0 108610 2 AC10572 5 100.0 121212 9 HS135L2	5 100.0 101290 9 AC00924	100.0 100685 8 AC02375	5 100.0 91240 2 AC10614	100.0 50210 9 AC07926	3523		Query Core Match Length DB I
dly,B.M. and Gurney,A.L. AN-2002;	bp DNA linear PAT 20-APR-200 342220.	LIGNMENTS	999	s mus ndosc	AL500526 Human	5151 Mycobact	18173 Dros 187 Mycoba	AE013309 Methan	095056 Leptosp	AL684904 Penicili AF422179 Arabidops	AL685230 Penici	AY038114 Mont	009519 Homo sap 038113 Montanoa	M27258 Human mut	AC124474 Mus mus	AC108796 Mu AC117630 Mu	AC101658 Mus mus	AC115064 Mus mus	AC128590 Ra AC018846 Ho	AC099146 Rattus	AC022311 Homo sa	AC016036 Home	AC131121 Mus mu	AC1057 AL0317	ACCOUNTY HOMO SAP	AC023754 Arabidop AC121474 Rattus n	AC106140 Rattus n	AC079261 Homo AY007367 Lycop	ARIB3492 Sequence ACO24250 Homo sap		⊢.

FEATURES	TITLE	AUTHORS	REFERENCE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	LOCUS	AR183492	KENCET L
Location/Qualifiers	Agonist antibodies Datent: US 6342220-A 36 29-TAN-2002:	Adams, C.W., Carter, P.J., Fendly, B.M. and Gurney, A.L.	1 (bases 1 to 15)	Unclassified. '	Unknown.	Unknown.		AR183492.1 GI:20227461	AR183492	Sequence 36 from patent US 6342220.	AR183492 15 bp DNA linear PAT 20-APR-2002		

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Direct Submission
Submitted (28-FEB-2000) G
University School of Medi
MO 63108, USA
Inis sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by restriction digest.
                                                                                                                clone. It may be shorter because we only sequence clone sections once, or longer because in the entire between neighboring.
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Submitted (09-MAY-2001) Department of Genetics,
University, 4444 Forest Park Avenue, St. Louis,
On Nov 8, 2000 this sequence version replaced g
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Genome Res. 8 (11), 1097-1108
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                                                                                                                                                                                             Contact: sapiens@watson.wustl.edu
------ Summary Statistics
Center project name: H_NH0684N02
                                                                                                                                                                                                                                                      Center: Wash:
Center code:
                                                                                                                                                                                                                                     Web site: http://genome.wustl.edu/gsc
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Primates;
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ed. No. 2.6e+02;
Mismatches 0;
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Catarrhini;
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9847074
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Ozersky, P., Abbott, A., Maupin, R. and The sequence of Homo sapiens BAC c. Unpublished
3 (bases 1 to 50210)
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Toward a
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VECTOR: pBACe3.6
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The clone sequenced to the left is RP11-943F15; the clone sequenced to the right is RP11-489013. Actual start of this clone is at base position 17386 of RP11-943F15; actual end is at base position 71706 of RP11-489013.
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The position of this clone was established as part of a collaboration between the Human Chromosome Y Mapping Project (Tomoko Kawaguchi, Helen Skaletsky, Laura G. Brown, Steve Roz and David C. Page at the Whitehead Institute for Biomedical Research, Cambridge MA) and the Washington University Genome segmencing Center, St. Louis MO.
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(ston, J.E. and Waterston, R.
ard a complete human genome)
nome Res. 8 (11), 1097-1108
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piens BAC clone RP11-1325K3
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8789 c 10023 q
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/db_xref="taxon:9606"
/chromosome="Y"
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/clone_lib="RPCI-11"
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(1998)
                                  and Ames, M. clone RP11-1325K3
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Submitted (09-MAY-2001) Department of Genetics,
University, 4444 Forest Park Avenue, St. Louis,
On Oct 26, 2000 this sequence version replaced composition of the composition of Genome Center of Composition of Genome Center of Composition of Co
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Submitted (26-OCT-2000) Genome
University School of Medicine,
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Direct Submission
Submitted (25-AUG-2000) Genome
University School of Medicine,
MO 63108, USA
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  Web site: http://genome.wustl.edu/gsc
Contact: sapiens@watson.wustl.edu
----- Summary Statistics
Center project name: H_NH1325K03
                                                                                                                                                                                           Center: Washington University Center code: WUGSC
Center project name:
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clone. It may be shorter because we or clone sections once, or longer because between neighboring data submissions. NOTICE: This sequence may not represent we only we provide a the entire insert of this y sequence overlapping e provide a small overlap

all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by restriction digest.

collaboration between the Human Chromosome Y Mapping Project (Tomoko Kawaguchi, Helen Skaletsky, Laura G. Brown, Steve Roand David C. Page at the Whitehead Institute for Biomedical Research, Cambridge MA) and the Washington University Genome Sequencing Center, St. Louis MO. PING INFORMATION:

position of this clone was established as position between the Human Chromosome Y Jaura G. Rozen,

SOURCE INFORMATION:

The RPCI-11 human BAC library was made from the blood of one male donor, as described by Osoegawa, K., Woon, P.Y., Zhao, B., Frengen, E., Tateno, M., Catanese, J.J. and de Jong, P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from Research Genetics, Inc. (http://www.resgen.com) or Pieter de Jong and coworkers at the Roswell Park Cancer Institute (http://bacpac.med.buffalo.edu) The clone sequenced to the left is RP11-1285C3; the clone to the right is RP11-943F15. Actual start of this clone i position 1 of RP11-1325K3; actual end is at base position RP11-1325K3. NEIGHBORING SEQUENCE VECTOR: pBACe3.6 INFORMATION: e sequenced is at base n 50210 of

The entire clone of H_NH1325K03 from 1 to 50210, is a 125 bp un tandem repeat. There are low quality areas, single clone areas, single chemistry areas and high quality mismatches through this clone. Digest cannot confirm any data within this clone do to the size of the tandem and its repeatative nature. Due to the tandem the exact overlap cannot be determined, but RP11-1325K3 RP11-943F15 are thought to overlap. The two clones are going the butt joined to give the best available representation of the region. throughout the

Location/Qualifiers

FEATURES

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BASE COUNT
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Prins,T.W., Spassova,M.I., Prins,M., Klein Lan Folkertsma,R.T., Hille,J. and Goldbach,R.W. Direct Submission Submitted (22-AUG-2000) Department of Virolgy, University, Binnenhaven 11, Wageningen 6709 PD Location/Qualifiers
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Eruction of a bacterial artificial chromosome (BAC) persicon esculentum cv. Stevens and its application ically map the Sw-5 locus

Breed. 5, 197-207 (1999)
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.W., Spassova.M
                                                                   /map="tightly linked to RFLP mark
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/gene="Sw5-c"
<23512. .>27327
/gene="Sw5-c"
/product="tospovirus resistance p
23512. .27327
/gene="Sw5-c"
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/chromosome="Y"
/map="Y"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="RP11-1325K3"
/clone_lib="RPCI-11"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; yta; Magnoliophyta; eudicotyledons; core eudicots; euasterids I; Solanales; Solanaceae; Solanum;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GI:15418711
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72 c 11695 g
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Pred. No.
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2e+02;
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                                                                                                    protein
                                                         protein"
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, The Netherlands
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pin C (Sw5-c),
rirus resistance
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GAIAGDVLYVIQKLLLSSIKKDDNSKISLCSIQIWEKTKDLKAQVETYYKSLKFTPSQ
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YLYATEINGYVDYEKLESLETRIQLMADNVGQLCTAFIVFEDEDEDEIVDIVENKP
PYLLGLIVLVEKEMKKIFLNELKASKFTQSKTFKDKKLPKGFSHHLNSLLVYLRNTKL QLQILDLSFRLSHMVSGLQLPSNLKMLVLREADLEKLTSFVAGLPSLEYLQLTNMYFH QSEWCLGDITFHKLKLLKLVQLDISRWDVSEESFPLLETIIIRWCLNLEEIPISFADV PTLKQIKLIHCTKKSLDDSAVRIKKDVEENEGNDRIDLIIKVSRNKLLCFVFEYLPAS YVTEINGYVNHEKLECLETRMHFMAYNVGQLCFAILGYDVSNFVDDGDKDDDDEEDED EDGDGDEDDILNKPPYLLCLIILVELEMKKIFLGEQKASKFTQSRTFKDKKLPKGFSH HLHSLLMYLRNKKLEKFPNNNSPQNIDIAIEFLLVFLDADVANHVINGNWLNEVMEKV /9777...41502
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AUTHORS
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Mizny, D.M., Adams, C., Adio-Oddola, B., Ali-osman, F.R., Allen, C., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbrooks, S.L., Amaratunge, K., Blankenburg, K., Bonnin, D., Bouck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Bulbay, C., Burch, P., Burkett, C., Burcell, K.L., Bydd, N.C., Chen, R., Chen, R., Chen, R., Chacko, J., Chavez, D., Cheveland, C.D., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., Davila, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A., Detlaney, K.R., Delgado, O., Denh, A.L., Ding, Y., Dinh, H.H., Douthwaite, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J., Detlaney, K.R., Delgado, O., Denh, A.L., Ding, Y., Dinh, H.H., Duyis, R., Garo, J., Garcia, A., Garner, T., Garza, N., Gill, R., C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Garlis, T., Garza, N., Garner, T., Garza, N., Gill, R., C., Earnhart, C., Harris, K., Hart, M., Havlak, P., Hawes, A., Hernandez, J., Harris, C., Harris, K., Hart, M., Havlak, P., Hawes, A., Hernandez, J., Hernandez, J., Hosson, E., Kelly, S., Huber, J., Hullyk, S., Hume, J., Jackson, L.E., Jacobson, B., Jia, Y., Johnson, R., Jolivet, S., Joulah, S., Kratovic, J., Kureshi, A., Landry, N., Leal, B., Lewis, L.C., Lewis, L., Li, J., Li, X., Lucier, A., Lucier, R., Luna, R., Ma, J., Loulseged, H., Lozado, R.J., Lu, X., Lucier, A., Lucier, R., Luna, R., Ma, J., Lu, X., Lucier, A., Lucier, R., Martindale, A., Martinez, E., Mosser, M., Mellor, J., Metzker, M., Mohabbat, K., Morgan, M., Morris, S., Mosser, M., Newtson, J., Newtson, N., Nguyen, A., Nguyen, N., Nguyen, N.
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norvegicus clone CH230-230F16, unordered pieces.
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KKVFQKEDCPPELQDVSQVVAEKCKGLPLVIVLVAGIIKKRKMEESWWNELKDSLFDY
LDCHSEQYSRATMQLSFDNLADCLKPCLLYMGMFPEDACIKVSALLSLWIAEDFVQNI
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QFQSFVWKQSRVSFSLSEENSKFASKTRKCSHQPLRSLITNGASFDIISLSSWIHNLR
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MLRGCFWEMEQLRHVEISDAEFDKQGLFEGSSKLENLRILKNIVGFPIDRADVLSRRC
PNLQQLHIEFSIDEAADSFCLTLENLTQLQKLRLSTECHSVLSRLQLPSNLKKLVLSN
TCIENLTSFLEGLPSLEYLQLRDIYFPHQSKEWCLGDITFHKLKLLKLVQLDISRWDV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEESFPLLETIIIRRCQNLEEIPISFADVPTLKQIKLIHCKKKSLDDSAVRIKKDVED
NEGNDRIDLIMKVSRNKLLCFVFEYLPASNLTPIIQK"
9668 c 9869 g 20265 t
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EKTKDLKAQVETYYKSLIFTPSQFPTFGGLTFMDSLLRKLNEMSRSKYGLDFLMKPIF
CNLEKEQSTLISNLEKELSSLSSIFRDVTKVHHEHKDLQRRTINLAYEAGVSIDSICA
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FDVRAWCIVSQTYNRRKLLQEILSQVTGSKDKGYEDDILADELRKSLMGKRYLIVLDD
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KASKFTQSKTFKDKKLPKGFSHHLNSLLVYLRDTKLKNFPNNVSAKNIGLSIELLLFF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KYNVFWHCFCSLPTILEEIKQINAEVTEMWSADIPLNPHYVAAPFKHLPTRHSNPITD
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Eutheria;
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Rodentia;
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Sciurognathi; Muridae; Murinae;
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2e+02;
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2 (bases I to 91240)
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5 Worley, K.C.
Direct Submission
5 Lobmitted (13-JUL-2002) Human Genome Sequencing Center, Departmen of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
5 Direct Submission
5 Direct Submission
6 Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
6 On Jul 11, 2002 this Sequence version replaced gi:18138661.
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* NOTE: This is a 'working draft
* consists of 54 contigs. The tr
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Web site: http://www.hgsc.bcm
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Direct Submission

L Submitted (09-MAR-2000) DNA Sequencing and Technology Center Stanford University, 855 California Avenue
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L Unpublished

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Direct Submission

Submitted (17-FEB-2000) DNA Sequencing and Technology Center, Stanford University, 855 California Avenue, Palo Alto, CA 94304, USA

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                                                                                                                                     Howng, B., I
Liu, S.,
Hansen.N.F.,
J., Theologis,
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), CA 94304,
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, CA 94304,
                                                                                                          Ecker, J.,
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Kim,C
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Direct Submission

Submitted (13-SEP-2000) DNA Sequencing and Technology Center,

Stanford University, 855 California Avenue, Palo Alto, CA 94304,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene prediction (Informatics
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ive', '-like' or 'similar to'.

irity but no significant protein similarity are described as motated based only on company that are annotated based only on company the software are described as 'hypothetical proteins'.

prediction programs used to predict genes include: Gramatics Group, Oak Ridge National Laboratory, Chris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ard, et al., CBS, Technical University of k,http://www.cbs.dtu.dk/NetPlantGene.html).
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         with similarity to
ive', '-like' or 's
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                43756. .46560
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                                                                                                                                                                               complement(join(46848.
47773. .47875,47963. .
                                                                                                                                                                                                                                                                                                                                                          SSVSSNGSVLYCLEITKNYHDSDSEIVDQEVEILMKKLNFIPTSVFTTDLQYVDFLDR
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43756
                                                                                                                                                                                                                                                                                             complement(46848.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="Similar to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /gene="F1B16.
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) F10A5,
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.48066,48156. .
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.48185))
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- 56367
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'gene="F1B16.
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Alsbrooks,S.L., Amaratunge,H.C., Are,J.R., Ayele,M., Banks,T.,
Barbaria,J., Benton,J., Binage,K., Blankenburg,K., Bonnin,D.,
Bouck,J., Bowie,S., Brieva,M., Brown,E., Brown,M., Bryant,N.P.,
Bouck,J., Bowie,S., Brieva,M., Brown,E., Brown,M., Bryant,N.P.,
Buhay,C., Burch,P., Burkett,C., Burrell,K.L., Byrd,N.C.,
Carron,T.F., Carter,M., Cavazos,S.R., Chacko,J., Chavez,D.,
Chen,G., Chen,R., Chen,Z., Chowdhry,I., Christopoulos,C.,
Cleveland,C.D., Cox,C., Coyle,M.D., Dathorne,S.R., David,R.,
Davila,M.L., Davis,C., Davy-Carroll,L., Dederich,D.A.,
Davila,M.L., Ding,Y., Dinh,H.H.,
Douthwaite,K.J., Davards,C.C., Elhaj,C., Escotto,M.,
Garner,T., Garza,N., Guraratne,P., Hale,S., Hamilton,K.J.,
Gaorell,J.H., Guevara,W., Gunaratne,P., Hale,S., Hamilton,K.,
Harris,C., Harris,K., Hart,M., Havlak,P., Hawes,A., Hernandez,J.,
Homsi,F., Howard,S., Huber,J., Holloway,C., Hollins,B.,
Jacobson,B., Jia,Y., Johnson,R., Jolivet,S., Joudah,S.,
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Karlsson,E., Kelly,S., Khan,U., King,L., Korvah,J., Kovar,C.,
Karlsson,E., Kelly,S., Khan,U., King,L., Korvah,J., Kovar,C.,
Karlsson,E., Lichtarge,O., Lieu,C., Liu,J., Liu,W., Loulseged,H.,
Maheshwari,M., Mapua,P., Martin,R., Martindale,A., Martinez,E.,
Massey,E., Mawhiney,E., McLeod,M.P., Meador,M., Morgan,M., Morris,S.,
Ngoger,M., Neil,D., Newtson,N., Nguyen,A., Nguyen,N.,
Oyledo,R., Pace,A., Paven,B., Dearv.J., Perez.L.
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Nguyen, N., Nickerson, E., Nwokenkwo, S., Oguh, M., Okwuonu, G., Oragunye, N., Oviedo, R., Pace, A., Payton, B., Peery, J., Perez, L., Peters, L., Pickens, R., Primus, E., Pu, L.L., Quiles, M., Ren, Y., Rives, M., Rojas, A., Rojubokan, I., Rolfe, M., Ruiz, S., Savery, G., Scherer, S., Scott, G., Shen, H., Shooshtari, N., Sisson, I., Sodergren, E., Sonaike, T., Sparks, A., Stanley, H., Stone, H., Sutton, A., Svatek, A., Tabor, P., Tamerisa, A., Tamerisa, K., Tang, H., Usmani, K., Vasquez, L., Vera, V., Villalon, D., Vinson, R., Wang, S., Ward-Moore, S., Warren, R., Washington, C., Watlington, S., Wu, C., Wu, Y., Wu, Y.F., Zhou, J., Zorrilla, S., Nelson, D., Direct Submission
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/codon_start=1
/protein_id="AAG13064.1"
/db_xref="GI:10120439"
/translation="MGSSTSGNCSSVSTTGLANSGSESDLRQRDLIDERKRKRKQSNRESARRSRMRKQKHLDDLTAQVTHLRKENAQIVAGIAVTTQHYVTIEAENDILRAQVLE
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*** SEQUENCING IN PROGRESS ***,
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Direct Submission
Submitted (18-MAY-2002) Human Genome Sequencing Cer
of Molecular and Human Genetics, Baylor College of
Baylor Plaza, Houston, TX 77030, USA
3 (bases 1 to 101282)
Worley, K.C.
Direct Submission
Submitted (24-JUL-2002) Human Genome Sequencing Cer
of Molecular and Human Genetics, Baylor College of
Baylor Plaza, Houston, TX 77030, USA
On Jul 19, 2002 this sequence version replaced gi:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.NOTE: This is a 'working draft' sequence. It currently consists of 62 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Center project name: GYVV
Center clone name: CH230-88M5
Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye:
Assembly program: Phrap; version 0.
Consensus quality: 53833 bases at 1.
Consensus quality: 58211 bases at 1.
Consensus quality: 60723 bases at 1.
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Center: Baylor College of Medicine
Center code: BCM
Web site: http://www.hgsc.bcm.tmc.ed
Contact: hgsc-help@bcm.tmc.edu

Contact: Project Information
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3 (bases 1 to 101290)
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Direct Submission
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Waterston, R.
Direct Submission
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Submitted, 4444 Forest Park Avenue, St. Louis, Missouri 63108,
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Center: Washington University Genome Sequencing Center code: WUGSC Web site: http://genome.wustl.edu/gsc Contact: sapiens@watson.wustl.edu Center project name: H_NH0489013
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between neighboring clone. It may be she clone sections once, This sequence may not represent ty be shorter because we only sonce, or longer because we data submissions. ent the entire only sequence of we provide a re insert of this e overlapping a small overlap

rnis sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >: 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by restriction digest.

MAPPING The posi INFORMATION: tion of this

position this clone was es tablished as pa rt of

> (Tomoko Kawaguchi, Helen Skaletsky, Laura G. Brown, Steve Ro; and David C. Page at the Whitehead Institute for Biomedical Research, Cambridge MA) and the Washington University Genome Sequencing Center, St. Louis MO. collaboration between the Human Chromosome Y Mapping Project Rozen,

NEIGHBORING The clone se Tateno, M., approach f libraries. SOURCE IN The RPCIand Research (http://bacpac.med.buffalo.edu)
VECTOR: pBACe3.6 donor, as COWOI INFORMATION: -11 human BAC library was made from the blood of one male s described by Osoegawa, K., Woon, P.Y., Zhao, B., Frengen, E., Catanese, J.J. and de Jong, P.J. (1998) An improved for construction of bacterial artificial chromosome s. Genomics 51:1-8. The clone may be obtained either from Genetics, Inc. (http://www.resgen.com) or Pieter de Jong ckers at the Roswell Park Cancer Institute SEQUENCE INFORMATION:

tandem re was position missing used sequence H_NH0489013 from data. epeat at bas 73076 to make the assembly however, equenced to the left is RP11-684N2; the clone sequenced t is RP11-329C15, 200 bp overlap. Actual start of this base position 1 of RP11-489O13; actual end is at base 076 of RP11-329C15. the tandem repeat base positon 1 to 32763 contains into RP11-684N2. The sequence cannot be guaranteed. Digest dai however, it still suggests there data ۵v

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              such
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Mammalia; E
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (13-MAR-2000) Sanger Centre, Hinxton, Cambridgeshire CB10 1SA, UK. E-mail enquiries: humquery@sanger.ac.uk Clone requests: clonerequest@sanger.ac.uk
On Feb 11, 2000 this sequence version replaced gi:6911919.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 annotated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 corresponding to the overlapping clone, only a small overlap as described above.
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2976. .4675
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4748. .4900
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899. .1064
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1. .121212
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5530. .6001
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8425. .8721
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13175. .13601
/note="MER39
14084. .14389
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/note="LTR23 repeat: 1

12332. .12630

/note="AluSp repeat: 1

12647. .12953
                    /note="L1PA14 repeat: matches 5590.
28538. .30098
                                                        /note="MER58A repeat: matches 1.
27578. .27688
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22083. .22510
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14477. .14767
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'note="MIR r
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18083
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6325. .16543
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sequencing vector: M13; 0%
Sequencing vector: plasmid; 100%
Chemistry: Dye-primer ET; 0% of reads
Chemistry: Dye-terminator Big Dye; 100% of reads
Assembly program: Phrap; version 0.990319
Consensus quality: 121880 bases at least Q40
Consensus quality: 123345 bases at least Q30
Consensus quality: 124257 bases at least Q20
Insert size: 188000; agarose-fp
Insert size: 126600; sum-of-contigs
Quality coverage: 4.12 in Q20 bases; agarose-fp
Quality coverage: 4.82 in Q20 bases; sum-of-contigs
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McPherson, J.D. and Wat
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------ Project Information --
project name: M_BB0550I15
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1. .127400
35507. .64851
/note="assembly_name:Contig13"
64952. .113805
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/note="assembly_name:Contigl1"
20789. .35406
/note="assembly_name:Contigl2"
                                                                                                                                                                                      /organism="Mus musculus"
/db_xref="taxon:10090"
/chromosome="UNK"
/clone="RP24-550I15"
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9974: gap of unknown length
20688: contig of 10714 bp in le
20788: gap of unknown length
35406: contig of 14618 bp in le
35506: gap of unknown length
64851: contig of 29345 bp in le
64951: gap of unknown length
113805: contig of 48854 bp in le
113905: gap of unknown length
115892: contig of 1987 bp in len
115992: gap of unknown length
117914: contig of 1922 bp in len
118014: gap of unknown length
122126: contig of 4112 bp in len
122226: gap of unknown length
127400: contig of 5174 bp in len
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E 4 (bases 1 to 129547)

S Cheuk, R., Shinn, P., Brooks, S., Buehler, E., Chao, Q.,
Johnson-Hopson, C., Khan, S., Kim, C., Altafi, H., Bei, B., Chin, C.,
Chiou, J., Choi, E., Conn, L., Conway, A., Gonzalez, A., Hansen, N.,
Howing, B., Koo, T., Lam, B., Lee, J., Lenz, C., Li, J., Liu, A., Liu, J.,
Liu, S., Mukharsky, N., Nguyen, M., Palm, C., Pham, P., Sakano, H.,
Schwartz, J., Southwick, A., Thaveri, A., Toriumi, M., Vaysberg, M.,
Yu, G., Davis, R., Federspiel, N., Theologis, A. and Ecker, J.
Direct Submission
Submitted (22-JUL-2000) Arabidopsis thaliana Genome Center,
Department of Biology, University of Pennsylvania, 38th and
Hamilton Walk, Philadelphia, PA 19104-6018, USA
                                                                                                                                                                                                                                                                                                                        Ecker, J.R.

Direct Submission

L Submitted (23-APR-1999) Arabidopsis thaliana Genome Center, Department of Biology, University of Pennsylvania, 38th Stree, Hamilton Walk, Philadelphia, Pennsylvania 19104-6018, USA (bases 1 to 129547)

Cheuk, R., Shinn, P., Brooks, S., Buehler, E., Chao, Q., Johnson-Hopson, C., Khan, S., Kim, C., Altafi, H., Bei, B., Chin, Chiou, J., Choi, E., Conn, L., Conway, A., Gonzalez, A., Hansen, N., Howing, B., Koo, T., Lam, B., Lee, J., Lenz, C., Li, J., Liu, A., Liu, S., Mukharsky, N., Nguyen, M., Palm, C., Pham, P., Sakano, H., Schwartz, J., Southwick, A., Thaveri, A., Toriumi, M., Vaysberg, Yu, G., Davis, R., Federspiel, N., Theologis, A. and Ecker, J. Direct Submission

L Submitted (28-JUN-2000) Arabidopsis thaliana Genome Center, Department of Biology, University of Pennsylvania, 38th and Hamilton Walk, Philadelphia, PA 19104-6018, USA

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Submitted (24-JAN-1999) Arabidopsis thaliana Genome Center,
Department of Biology, University of Pennsylvania, 38th Street
Hamilton Walk, Philadelphia, Pennsylvania 19104-6018, USA
2 (bases 1 to 129547)
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eu
Rosidae; eurosids II; Brassicales; Brassicaceae; Arak
1 (bases 1 to 129547)
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118015. .122126
/note="assembly_name:Contig8"
122227. .127400
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/note="assembly_name:Contig6"
115993. .117914
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38th and
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Vaysberg,M.,
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38th Str
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Yu,G.,
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/product="f10A5.3"
/product="f10A5.3"
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LERINVYYNEASGGRYVPRAVLMDLEPGTMDSIRSGPYGQIFRPDNFVFGQSGAGNNW
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PDRMMLTFSVFPSPKVSDTVVEPYNATLSVHQLVENADECMVLDNEALYDICFRTLKL
STPSFGDLNHLISATMSGVTCSLRFPGQLNSDLRKLAVNLIPFPRLHFFMVGFAPLTS
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complement(join(8344. .8471,8704. .8782,8967. .9293))
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/db_xref="GI:9369356"
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KQVIVINDMFPGPILNATANDIIVVKDQIGSYFYFPTLLLQKAAGGYGAIRIYPPELV
PVPFPKPDEEYDILIGDWFYLDHTVMRASLDAGHSLPNPDGILFNGRGPEETFFAFEP
GKTYRLRISNVGLKTCLNFRIQDHDMLLVETEGTYVQKRVYSSLDIHVGQSYSILVTA
KTDPVGIYRSYYIFATARFTDSYLGGIALIRYPGSPLDPVGQGPLAPALQDFGSSVEQ
ALSIRMDLNVGAARSNPQGSYHYGRINVTRTIILHNDVMLSSGKLRYTINGVSFVYPE
TPLKLVDHFQLNDTIIPGMFPVYPSNKTPTLGTSVVDJHYKDFIHIVFQNPLFGLESY
HIDGYNFFVVGYGFGAWSESKKAGYNLVDAVSRSTVQVYPYSWTAILIAMDNQGMWNV
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join(5713. .6109,6903. .7172,7261. .7937)
/note="putative tubulin beta-4 chain sp|P50262|TBB4_PORPU;
similar to ESTs dbj|AV54234.1, dbj|AV536568.1,
dbj|AV543492.1, dbj|AV542234.1, and gb|W43629.1"
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/translation="MSKKKKTQSLSMEDHRDGKGKKRDDFFDSIDYEISSILLQLSHPVIFSSDSPVLHKWGRTKKRSSSSSHLHPPEPVIKSLPCMADVGDMGSNSSSSCLTGEAKKTNSQIEKSEFFSPRGVHCYLFIFCFPFCIKKSEEGFQRVPNKVQVAAQARQRRLGL
                                                                                                         /product="F10A5.4"
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/db_xref="GI:9369385"
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3056. .3416,3520. .3885,3968. .4046,4126. .4335)
/note="similar to L-ascorbate oxidase homolog precursor
sp[Q00624|ASO_BRANA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /translation="MALPLPLIFLIFSHLFVSGVRSTSFIMVNKCEYTVWPGLLSNAGVPPLPTTGFVLQKGEERTISAPTSWGGRFWGRTQCSTDTDGKFTCLTGDCGSGTLECSGSGATPPATLAEF"
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                                                                                                                                                                                                                     /evidence=not_
                                                                                                                                                                                                                                                            /codon_start=1
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1. .129547
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SPSNTASLLNCEPLRVQPRKLKRCASHIYISRLIKVLQTSKSSLATLNQNEQRYSEMS
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YNHNVSQQQHRLMTAAAAAMSMSHQHNNPSRAVMNRQEHHFPLIYEDTRTPLQLLCNEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /translation="MAISKALIASLLISLLVLQLVQADVVRLFHHKLIILNIILMFSK
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complement(18355. . 18426)
/product="trna-Thr"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /translation="MKAATRPIHAVTTWVRRQPPKVKGFLGVVSAMTALVLLRVIVHD HDNLFVAAEAVHSLGISVLIYKLTKERTCAGLSLKSQELTALFLAVRLYCSFVMEFDI HTLLDSATLVTTLFVIYMIRFKLKASYMDDKDNFAIYYVVIPCVVLSVLIHPSTHHHI INKISWAFCVYLEAVSVLPQLRVMQNTKPFTAHYVFALGIARFLSCAHWVLQVLDTRG RLLTALGYGFWPIMVLLSEIVQTFILADFCYYYVKSLMGGQLVLRLPSGVV" join(17031. 17238,17321. 17502) /note="similar to GAST-like gene product gb|AAB97006.1; similar to ESTs dbj|AV536142.1, and dbj|AV566951.1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         join(10894. .11116,11285. .11484,11708
12074. .12198,12327. .12375)
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gb|R30294.1, dbj|AV545596.1, and gb|AI992442.1"
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/product="F10A5.5"
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gb[R30294.1,
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/note="hypothetical protein
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E (bases 1 to 145520)

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S Birren,B., Linton,L., Nusbaum,C., Lander,E., Allen,N., Anderson,M.,

Birren,B., Linton,L., Nusbaum,C., Lander,E., Allen,N., Anderson,M.,

Birren,B., Linton,L., Nusbaum,C., Lander,E., Boukhgalter,B.,

Baldwin,J., Barna,N., Beckerly,R., Boguslavkiy,L., Boukhgalter,B.,

Brown,A., Castle,A., Colangelo,M., Collins,S., Collymore,A.,

Cooke,P., DeArellano,K., Dewar,K., Domino,M., Donelan,L., Doyle,M.,

Ferreira,P., Cardyna,S., Grant,G., Hagos,B., Heaford,A., Horton,L.,

Galagan,J., Gardyna,S., Grant,G., Hagos,B., Heaford,A., Horton,L.,

Howland,J.C., Johnson,R., Jones,C., Kann,L., Karatas,A., Klein,J.,

McEwan,P., McGurk,A., McKernan,K., McLaughlin,J., Marquis,N.,

Morrow,J., Naylor,J., Norman,C.H., O'Connor,T., O'Donnell,P.,

Peterson,K., Pollara,V., Riley,R., Roy,A., Santos,R., Severy,P.,

Stange-Thomann,N., Stojanovic,N., Subramanian,A., Talamas,J.,

Tesfaye,S., Tirrell,A., Vassiliev,H., Vo,A., Wheeler,J., Wu,X.,

Wyman,D., Ye,W.J., Zimmer,A. and Zody,M.
                                                  sequencing reads that have not been assembled into contigs. Runs of N are used to separate the reads and the order in which they appear is completely arbitrary. Low-pass sequence sampling is useful for identifying clones that may be gene-rich and allows overlap relationships among clones to be deduced. However, it should not be assumed that this clone
                                                                                                                                                                                                                                                           nitted (19-NOV-1999) Whitehead Institute/MIT Center for Genome earch, 320 Charles Street, Cambridge, MA 02141, USA Jul 13, 2000 this sequence version replaced gi:7108033. repeats were identified using RepeatMasker: t, A.F.A. & Green, P. (1996-1997)
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                          be sequenced to
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/product="F10A5.10"
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Direct Submission
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Waterston, R.
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Sulston, J.E. and Waterston, R
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                                                                                                                                                                                                                                                  NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.
MAPPING INFORMATION: Mapping information for McPherson, Department of
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Web site: http://genome.wustl.edu,
Contact: sapiens@watson.wustl.edu
Center project name: H_NH0146M24
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was provided by Dr. Jo. Washington University,
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additional information about the map position of see http://genome.wustl.edu/gsc
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The RPCI-11 human BAC library was made from the blood of one male donor, as described by Osoegawa, K., Woon, P.Y., Zhao, B., Frengen, E., Tateno, M., Catanese, J.J. and de Jong, P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from Research Genetics, Inc. (http://www.resgen.com) or Pieter de Jong and coworkers at the Roswell Park Cancer Institute
                                                                                 The
                                                                                                       (http://bacpac.med.buffalo.edu)
VECTOR: pBACe3.6
NEIGHBORING SEQUENCE INFORMATION
                                                    to the right
RP11-146M24.
                         position
                  clone sequenced to the left is RP11-457A20; the right is RP11-36C8. Actual start of this ition 1 of RP11-146M24; actual end is at base
                                                                                                                                                                                                                                                                                                                                                                                                            NFORMATION:
                                                                                                                   INFORMATION:
                                                      the clone is
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at base
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There are the clone polymorphic base pair diff RP11-146M24 and RP11-36C8. Location/Qualifiers differences in the overlap between

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143. .213
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1583. .1629
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3328. .3532
/rpt_family="MIR"
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                                                             Direct Submission

L Submitted (08-AUG-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA

Center: Baylor College of Medicine
Center code: BCM
Web site. L.
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          Contact: hgsc-help@bcm.tmc.edu
------ Project Information
Center project name: KATP
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Agonist antibody; thrombopoietin receptor; TPO-R; thrombopoietin; DIC; megakaryocyte; platelet; immunological; hematopoietic; thrombocytopenia; bone marrow hypoplasia; disseminated intravascular coagulation; anemia; myelodysplasia; myelotoxic chemotherapy; leukaemia; tumour; MuSK; CDR; neuromuscular; muscular dystrophy; complementarity determining region; variable heavy chain; variable light chain. vu. vi.

RESULT 1
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                                                                                                        2780 A;
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                                                                                                        2460 C;
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Pred. No.
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                                  Score 14;
Pred. No.
0; Mismato
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4.5e+02;
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hes 0;
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RESULT 6
AAC56643/c
ID AAC56643:
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AC AAC56643;
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DT 25-JAN-20:
XX
DE Eucalyptu:
XX
KW 'Plant; tra
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ABL19870
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                                                                                                                                                  Дb
                                                                                                                                                                                             Query Match
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Matches 14
                                                                                                                                                                                                                                                             The invention relates to an isolated nucleic acid detection reagent capable of detecting 1000 or more genes from Drosophila. The invention useful in developmental biology and in elucidating cell signalling and cell-cell interactions in higher eukaryotes for the development of insecticides, therapeutics and pharmaceutical drugs. The invention discloses genomic DNA sequences (ABL16176-ABL30511), expressed DNA sequences (ABL01840-ABL16175) and the encoded proteins (ABB57737-ABB72072).

The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPG at ftp.wipo.int/pub/published_pct_sequences.
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interactions -
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11-JUL-2000;
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                                                                    AAC56643;
                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                        Claim
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                        Eucalyptus
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                                              JAN-2001
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Similarity 100.0%;
14; Conservative
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                       grandis
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000US-0614150.
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                                                                                                                                                                                                                                            2780 A;
                                                                                           DNA;
                       transcription factor DNA sequence #514.
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                                                                                                                                                   4064
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                                                                                                                                                                                                                                                                                                                                                                                                                                       detection reagent for detecting 1000 or for elucidating cell signalling and cell
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                                                                                           B₽.
                                                                                                                                                                                              Score 14; DB 23;
Pred. No. 4.7e+02;
0; Mismatches 0;
gene
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                                                                                                                                                                                                                                           2777 T;
                                                                                                                                                                                                                    Length 10426;
                                                                                                                                                                                                Indels
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transcr

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factor;

expression;

eucalyptus; pine;

acacia;

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RESULT 7
AAC55981/c
ID AAC55981;
XX
AC AAC55981;
XX
DT 25-JAN-2001 (first
XX
DE Eucalyptus grandis t
XX
KW Plant; transcription
KW poplar; sweetgum; te
KW basic helix-loop-hel
KW homeodomain zipper;
KW type 2 Cys2His2; CCA
XX
OS Eucalyptus grandis.
XX
PN WO200053724-A2.
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Matches 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 woody plant e.g. a eucalyptus, pine, acacia, poplar, sweetgum, teak, or mahogany species or to modify the activity of a polypeptide in a plant. The transcription factors of the present invention are members from the following families of regulatory proteins: bZIP, bZIP family of G-box binding factors, basic helix-loop-helix zipper, homeotic/homeodomain/homeobox/MADS, homeodomain zipper, LIM domain, AP, and EREBs, zinc finger domains of type 2 Cys2His2, CCAAT box elements
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to novel plant transcription factors from Eucalyptus grandis or Pinus radiata. The present sequence is the coding sequence for one such transcription factor. The transcription factor may be used to produce a plant having modified gene expression such as a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated polynucleotide encoding a producing a plant e.g. a woody plant, pl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-SEP-2000
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18-AUG-1999;
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                                                                                                                                      Plant; transcription factor; gene expression; eucalyptus; pine; poplar; sweetgum; teak; mahogany; bZIP; G-box binding factor; basic helix-loop-helix zipper; homeotic; homeodomain; homeobox; homeodomain zipper; LIM domain; AP2; EREBs; zinc finger domain;
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                                                                                                             homeodomain zipper; LI type 2 Cys2His2; CCAAT
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MYB.
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14; Conservative
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99US-0149485
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R CHALLENGE FORESTS LTD.
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                                                                                                                element;
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smatches 1;
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nt, preferably eucalyptus or pine,
modified activity of a polypeptid
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05-MAR-1999;
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Pred. No. 8.4e
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RESULT 10
ABA62712/c
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AC ABA62712;
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DE Human foe
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OS Homo sapi
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PN WO2001572
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PD 09-AUG-20
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PF 30-JAN-20
PR 04-FEB-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the tissues they were isolated from human, mouse, chicken and rat chemotactic; proliferative; immunomodulatory; haematopoietic; cytostatic; antibacterial; antifungal; antiviral; antidiabetic; anticonvulsant; and antidepressant. The serent feetive:

anticonvulsant; and antidepressant. The serent feetive:

identification:
                                                                   30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      anticonvulsant; and antidepressant. The SESTs can be used for gene therapy and in vaccines. The SESTs are useful as probes for the identification and isolation of full-length cDNAs and genomic DNA molecules which correspond to the SESTs. Proteins encoded by the SESTs are useful in assays for determining biological activity and raising antibodies. They may be useful for treatment of autoimmune disorders (multiple sclerosis, insulin dependent diabetes), allergic conditions (asthma), myeloid or lymphoid cell deficiencies, wounds, burns, ulcers, osteoporosis, osteoarthritis, central nervous system disorders (Alzheimer's, Parkinson's, Huntington's disease, stroke), coagulation disorders (haemophilia, thrombosis), inflammatory disorders (Crohn's disease), tumours, bacterial, fungal or viral infections, depression and psoriasis. AAA45936 to AAA45931 represent linker variants which are given in the exemplification of the present invention.
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RESULT 11
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ID ABA30013:
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AC ABA30013:
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DT 23-JAN-20
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PN W02001572
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PD 09-AUG-20
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PF 30-JAN-20
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PR 26-MAY-20
PR 27-SEP-20
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Note: The sequence data for this patent did not form printed specification, but was obtained in electronic from WIPO at ftp.wipo.int/pub/published_pct_sequences
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2000US-0207456.

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2000US-0632366.

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2000GB-0024263.
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    ention provides a number of single exon nucleic acid e derived from genomic sequences expressed in the human be used to measure gene expression in brain cell samples, e the diagnosis and improved treatment of nervous system
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                                                                                                                                                                                                                                                 l DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pressed exon; gene expression analysis; probe;
heimer's disease; multiple sclerosis; schizophrenia;
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93.3%;
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RESULT 13
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ID AAK36915;
XX
AC AAK36915;
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AC AAK36915;
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DT 06-NOV-20
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DE Human bon
KW Human; bo
KW Microarra
XX
OS Homo sapi
XX
PN WO2001572
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PN WO2001572
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PF 30-JAN-20
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27-SEP-2000;
04-OCT-2000;
                                                                                   probes which are derived from bone marrow. They can be used samples, which may enable the such as lymphoma, leukaemia ar the probes of the invention.
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epilepsy a
invention.
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2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-0236359.
2000US-0236359.
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RESULT 14
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ID AAI17764;
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AC AAI17764;
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DT 12-OCT-20
XX
Probe #76
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PPO 12-OCT-20
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OS HOMO Sapi
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OS AUG-20
PR 26-MAY-20
PR 26-MAY-20
PR 21-SEP-20
PR 26-MAY-20
PR 26-MA
RESULT 15
AAI42716/c
ID AAI42716;
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AC AAI42716;
XX
DT 17-OCT-20;
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DE Probe #11;
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21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
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Note: The sequence data for this patent did not form part of the printe specification, but was obtained in electronic format directly from WIPC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Note: The sequence data for specification, but was obtain at ftp.wipo.int/pub/published
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                                                                                                                                   The present invention relates to single exon nucleic acid probes (SENP). The present sequence is one such probe. The probes are useful for producing a microarray for predicting, measuring and displaying gene expression in samples derived from human placenta. The probes are useful for antenatal diagnosis of human genetic disorders.
                                                                                                                                                                                                                  Human genome-derived single exon nucleic acid probes analyzing gene expression in human placenta -
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US-09-694-531-
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                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/694, CURRENT FILING DATE: 2000-10-23 PRIOR APPLICATION NUMBER: 09/198,955 PRIOR APPLICATION NUMBER: 1343/97 PRIOR APPLICATION NUMBER: 1343/97 PRIOR APPLICATION NUMBER: 1344/97 PRIOR APPLICATION NUMBER: 1344/97 PRIOR APPLICATION NUMBER: 60/067,249 PRIOR APPLICATION NUMBER: 60/067,249 PRIOR APPLICATION NUMBER: 60/067,249 PRIOR APPLICATION NUMBER: 60/067,240 PRIOR APPLICATION NUMBER: 60/067,240 PRIOR APPLICATION NUMBER: 09/073,684 PRIOR APPLICATION NUMBER: 09/073,684 PRIOR APPLICATION NUMBER: 09/184,217 PRIOR EILING DATE: 1998-05-06
     Query Match
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Matches 14
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NUMBER OF SE
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                                                                        TYPE:
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LENGTH: 10
TYPE: DNA
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OR APPLICATION NUMBER: 60/067,240
OR FILING DATE: 1997-12-02
OR APPLICATION NUMBER: 09/073,684
OR FILING DATE: 1998-05-06
OR APPLICATION NUMBER: 09/184,217
OR FILING DATE: 1998-11-02
                                                                                                                                                                                                                                                                                                                                                          ICANT: Kongsbak, Lars E OF INVENTION: No. 636. REFERENCE: 5378.200-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            No.
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APPLICATION NUMBER: 60/067,249
PTITUG DATE: 1997-12-02
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Bjornvad, Mads
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NAME: WILLIAMS, JR. JOSEPH A.
REGISTRATION NUMBER: 38,659
REFERENCE/DOCKET NUMBER: 2786
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-474-0448
                                                                                                                                                                                                                                                                                                                                                                                                                                     US/08/656,984A

LATION: 435

APPLICATION DATA:
APPLICATION NUMBER: US 07/827,689

FILING DATE: 27-JAN-1992

PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 26

PRIOR APPLICATE: 26
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FILING DATE: 18-MAY-19
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 07-JUN-19
                                                       -656-984A-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/656,984A
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APPLICANT: Kilgannon, Patrick D.
TITLE OF INVENTION: ICAM-4 Materials and Methods
NUMBER OF SEQUENCES: 42
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APPLICATION
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APPLICATION
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ZIP: 60606-
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07-JUN-1995
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05-AUG-1993
                                                                                                                                                                                                           AMS, JR. JOSEPH A. NUMBER: 38,659
CKET NUMBER: 2786
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18-MAY-1994
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22-JAN-1993
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05-JUN-1992
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                                                                                                                                   Patent No. 6057435
GENERAL INFORMATION:
APPLICANT: Godowsk
APPLICANT: Gurney,
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Dreger, Ginger R.
REGISTRATION NUMBER: 33,055
REFERENCE/DOCKET NUMBER: P113
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-3216
TELEPHONE: 650/952-9881
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 3355 base pairs
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/933,821
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Godowski, Paul J. APPLICANT: Gurney, Austin L. TITLE OF INVENTION: Tie Ligal NUMBER OF SEQUENCES: 17
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COMPUTER: IBM F
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech
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COMPUTER READABLE MEDIUM TYPE: 3.
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APPLICANT: Gurney, Austin L
TITLE OF INVENTION: Tie Liga
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                    ADDRESSEE:
STREET: 1
CITY: Sout
STATE: Cal
COUNTRY: U
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CITY: South San F:
STATE: California
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STRANDEDNESS: Single
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FORM: .5 inch,
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                                 US-09-136-828-
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Patent No
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Query Match
Best Local
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TELEPHONE: 650/225-3216
TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 3355 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMAT
                                                                                                    TELEFAX: 6
                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Ginger R.
REGISTRATION NUMBER: 33,055
REFERENCE/DOCKET NUMBER: P113
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-3216
TELEFAX: 650/952-9881
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FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INF
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                                                                              SEQUENCE (
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                                                                                                                                                                                                                               SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE
ADDRESSEE:
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SOFTWARE:
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                                                                                                                                                                                                                                                                                                                                      CITY:
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                                                                            N FOR SEQ ID NO:
CHARACTERISTICS:
: 3355 base pair
                                                      Nucleic Acid
DNESS: Single
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ilarity 93.3%;
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IBM PC compatible
YSTEM: PC-DOS/MS-DOS
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RESULT 8
US-09-332-928A-3
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US-09-136-801-
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/332,928A
FILING DATE: 14-Jun-199
CLASSIFICATION: <Unknown>
PRIOR APPLICATION NUMBER: 08/933,821
FILING DATE: <Unknown>
APPLICATION NUMBER: 08/933,821
FILING DATE: <Unknown>
APPLICATION NUMBER: 33,055
REGISTRATION NUMBER: 33,055
REGISTRATION NUMBER: 33,055
REFERENCE/DOCKET NUMBER: P1130
TELEPHONE: 650/952-9881
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 3355 base pairs
TYPE: Nucleic Acid
                                                                 Sequence 3, Application Patent No. 6413770
GENERAL INFORMATION:
APPLICANT: Godowsk
APPLICANT: Gurney,
APPLICANT: Hillan,
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GENERAL INFORMATION:
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Best Local
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   APPLICANT:
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ADDRESSEE: Genente
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TITLE OF INVENTION: Tie Ligands
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815 AGCCGTAACATGAAC 829
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Similarity 93.3%;
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CITY: South San
STATE: Californi
COUNTRY: USA
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STRANDEDNESS: Single
TOPOLOGY: Linear
Roy, Margaret
Ferrara, Napoleone
Tumas, Daniel
                                           Botstein, David
Goddard, Audrey
                                                                      Godowski, Paul J.
Gurney, Austin
Hillan, Kenneth
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Pred. No. 93;
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US-09-332-929-3
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Patent No. 6420542
GENERAL INFORMATION:
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Best Local
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TITLE OF INVENTION: '
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRES:
                            PRIOR APPLICATION I APPLICATION IFILING DATE:
                                                                                                               OPERATING SYSTEM: PC-D
SOFTWARE: WinPatin (Ge
CURRENT APPLICATION DATA:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/136,
                                                                                                                                                                                                                                                                                                     APPLICANT: Godowski, Paul J. APPLICANT: Gurney, Austin L. TITLE OF INVENTION: Tie Ligands NUMBER OF SEQUENCES: 17
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ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inc
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   ATTORNEY/AGENT NAME: Drege:
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ADDRESSEE: Gen-
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CLASSIFICATION:
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STATE: Cali
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lilarity 93.3%;
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SEQ ID NO: 3
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                                           TION DATA:
                                                                                                                                        : 3.5 inch, 1.44 Mb floppy disk IBM PC compatible YSTEM: PC-DOS/MS-DOS
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NUMBER: P1
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INFORMATION FOR SEQ
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REFERENCE/DOCKET NUMBER: P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-3216
TELEFAX: 650/952-9881
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LENGTH: 3355 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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                                                                                                                   R APPLICATION NUMBER: 60/052,/33
R FILING DATE: 1997-07-08
R FILING DATE: 1997-07-08
ER APPLICATION NUMBER: 60/052,795
ER FILING DATE: 1997-07-08
ER APPLICATION NUMBER: 60/051,91
ER APPLICATION NUMBER: 60/051,92
FTIING DATE: 1997-07-08
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R APPLICATION NUMBER: 60/052,73
R FILING DATE: 1997-07-08
R APPLICATION NUMBER: 60/05
R FILING DATE: 1997-08-18
R APPLICATION NUMBER: 60/05
R FILING DATE: 1997-08-18
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R FILING DATE: 1997-08-18
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APPLICATION NUMBER:
DATE: 1997-07
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APPLICATION NUMBER: 6
TWO DATE: 1997-07-
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No. 9;
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RESULT 12
US-08-840-466A-21
; Sequence 21, Appl:
; Patent No. 626156;
; Patent INFORMAS
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                                                                  COUNTRY: USA
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, 'CORRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match
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R APPLICATION NUMBER: 60/055,953
R FILING DATE: 1997-08-18
R APPLICATION NUMBER: 60/055,950
R FILING DATE: 1997-08-18
R APPLICATION NUMBER: 60/055,947
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 1997-09-12
APPLICATION NUMBER: 60/058,664
FILING DATE: 1997-09-12
APPLICATION NUMBER: 60/058,660
FILING DATE: 1997-09-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 1997-0 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           : Patent
129
477
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CITY: Was
STATE: D.
COUNTRY:
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O'Brien, Alison D.
Wachtel, Marian R.
INVENTION: Method Of Stimulating An Immune Response
By Administration Of Host Organisms That
Alone Or As A Fusion Protein With One Or
Antigens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATION:
  SIFICATION:
                    ICATION NUMBER: US/08/840,466A
NG DATE: 18-Apr-1997
                                                                                                                                                                                                                                                                                                                      ESSEE: Finnegan, Henderson,
Dunner, L.L.P.
ET: 1300 I Street, N.W., Su
                                                                                                                                                                                                                                                                                                                                                                 DENCE ADDRESS:
ESSEE: Finnega
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ION NUMBER: 60/055,947
ATE: 1997-08-18
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STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
SEQUENCE DESCRIPTION: SEQ ID NO:
US-08-840-466A-21
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US-09-696-188B-21
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REGISTRATION NUMBER: 43,505
REFERENCE/DOCKET NUMBER: 04999
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4400
TELEFAX: (202) 408-4400
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
LENGTH: 3106 base pairs
TYPE: nucleic acid
STRANDFONESS: 510010
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Best Local
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vers
CURRENT APPLICATION NUMBER: US/09/696,188B
FILING DATE: 26-Oct-2000
CLASSIFICATION: <Unknown>
PRIOR APPLICATION NUMBER: 08/840,466
FILING DATE: 1997-04-18
AFTORNEY/AGENT INFORMATION:
NAME: Boone, Laural S.
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                                           SEQUENCE CHARACTERISTICS:
LENGTH: 3106 base pa
                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4000
TELEFAX: (202) 408-4400
MATION FOR SEQ ID NO: 21:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnega
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CITY: Washi
STATE: D.C.
COUNTRY: US
STRANDEDNESS:
TOPOLOGY: line
                                                                                                                                     NAME: Boone, Laural S. REGISTRATION NUMBER: 4
REFERENCE/DOCKET NUMBE
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O'Brien, A
Wachtel, I
                 nucleic acid
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1300 I Street,
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  Marian L.
n, Alison D.
l, Marian R.
l, Method Of Stimulating An Immune Response
N: Method Of Stimulating An Immune Response
By Administration Of Host Organisms That
Alone Or As A Fusion Protein With One Cr
              single
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                                                                                                                                    BER: 43,505
NUMBER: 049
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Pred. No.
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RESULT
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RESULT 15
US-09-696-188B-20
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US-09-696-188B-21
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GENERAL INFORMATION:
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sequence 20,
                                                                                                                                                                                                                                                              INFORMATION FOR SEQUENCE C
                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/840,466A
FILING DATE: 18-Apr-1997
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Boone, Laural S.
REGISTRATION NUMBER: 43,505
REFERENCE/DOCKET NUMBER: 04995.0029-00000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4000
TELEFAX: (202) 408-4400
FORMATION FOR SEQ ID NO: 20:
                                                                                                                         Local
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V
CURRENT APPLICATION NUMBER: US/08/840,466A
                                                                                                                                                                               MOLECULE
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow,
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CITY: Wa
STATE: I
COUNTRY:
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                                                                                                                                                                                                                     LENGTH: 3131 base pairs TYPE: nucleic acid STRANDEDNESS: single
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Mckee, Marian L.

O'Brien, Alison D.

Wachtel, Marian R.

FINVENTION: Method Of Stimulating An Immune Response

By Administration Of Host Organisms That Exp

Alone Or As A Fusion Protein With One Or Mo
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                                                                                                                                                                             TYPE: DNA (genomic)
DESCRIPTION: SEQ ID
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DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TRY: USA
20005-3315
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1300 I Street, N.W.,
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IMP PC compatible
COMPUTER: Patentin Release #1.0, Vers.

SOFTWARE: Patentin Release #1.0, Vers.

CURRENT APPLICATION NUMBER: US/09/696,188B
FILING DATE: 26-Oct-2000
CLASSIFICATION: <Unknown>
PRIOR APPLICATION NUMBER: 08/840,466
FILING DATE: 1997-04-18
ATTORNEY/AGENT INFORMATION:
NAME: BOONE, Laural S.
REGISTRATION NUMBER: 04995.0029-0.

TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4400
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 3131 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
SEQUENCE DESCRIPTION: SEQ ID NO: 20:
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Search completed: June Job time : 31.2 secs
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Best Local Similarity 100.0%;
Matches 13; Conservative
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GENERAL INFORMATION:
APPLICANT: Stew
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson
Dunner, L.L.P.
STREET: 1300 I Street, N.W., S
CITY: Washington
STATE: D.C.
COUNTRY: USA
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APPLICANT: Stewart, C. Near

Mckee, Marian L.

O'Brien, Alison D.

Wachtel, Marian R.

Wachtel, Marian R.

TITLE OF INVENTION: Method Of Stimulating An Immune Response

By Administration Of Host Organisms That Express Intimin

Alone Or As A Fusion Protein With One Or More Other

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2: /cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*

3: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*

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5: /cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*

6: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*

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US-09-910-943-412
US-10-072-152-7
US-09-938-842A-2733
US-09-948-783-79
US-09-892-877-115
US-09-948-783-115
US-09-948-783-115
US-10-025-524-33
US-09-897-306-2
US-10-028-072-277
US-10-121-049-277
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Sequence 5, Appli

Sequence 6, Appli

Sequence 412, App

Sequence 7, Appli

Sequence 2733, Appli

Sequence 80, Appl

Sequence 79, Appl

Sequence 115, App

Sequence 33, Appl

Sequence 27, Appli

Sequence 27, Appli

Sequence 277, App

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ALIGNMENTS

RESULT 1 US-09-864-761-8 Sequence Patent No

1-8479/ 479, Ap US2002

oplication US/09864761 20048763A1

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GENERAL INFORMATION:

APPLICANT: Penn, Sharron G.
APPLICANT: Hank, David R.
APPLICANT: Hank, David R.
APPLICANT: Hank, David R.
APPLICANT: Hensel, David R.
APPLICANT: Cheb, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
TITLE OF INVENTION: HUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/23,366
PRIOR APPLICATION NUMBER: US 60/23,366
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PRILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PRILING DATE: 2001-01-30
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                                                                                                                                       NAME/KEY: unsure
LOCATION: 532, 536,
OTHER INFORMATION: a
3-09-897-306-5
                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 5, Application US/09897306 Patent No. US20020123054A1
                                                                                                                                                                                                                                                                              APPLICANT: Patterson, Chandra APPLICANT: Murry, Lynn E. TITLE OF INVENTION: HUMAN ANGIOPOIETIN FILE REFERENCE: PC-0048 CIP CURRENT APPLICATION NUMBER: US/09/897,306 CURRENT FILING DATE: 2001-07-02 NUMBER OF SEQ ID NOS: 14
                                                                                                                                                                                                                                                                        SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Hillman, Jennifer L. APPLICANT: Gorgone, Gina A. APPLICANT: Patterson, Chandra
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OTHER INFORMATION:
09-864-761-8479
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                                                                                                                                                                                     NAME/KEY: misc_feat
OTHER INFORMATION:
                                                                                                                                                                                                               ORGANISM: HOMO FEATURE:
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OTHER
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LENGTH: 556
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N HELA, SIGNAL = 2.8
N ADULT LIVER, SIGNAL = 3.0
N BONE MARROW, SIGNAL = 3.0
N FETAL LIVER, SIGNAL = 3.0
N HEART, SIGNAL = 3.0
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FEATURE:

NAME/KEY: misc_feature:

LOCATION: (1)..(776);

OTHER INFORMATION: r

US-09-910-943-412
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; NAME/KEY: unsure
; LOCATION: 475, 506, 510
; OTHER INFORMATION: a, t
US-09-897-306-6
RESULT 5
US-10-072-152-
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SEQ ID NO 412
LENGTH: 776
                                                                                                      Matches
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APPLICANT: Hillman, Jennifer L.
APPLICANT: Gorgone, Gina A.
APPLICANT: Patterson, Chandra
                                                                                                                                                                                                                                                                                                                                                                               GENERAL
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Sequence 412, Application US/09910943
Patent No. US20020081610A1
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APPLICANT: Altman, Curtis
TITLE OF INVENTION: Assays and Materials for
FILE REFERENCE: 7529/1G148US1
CURRENT APPLICATION NUMBER: US/09/910,943
CURRENT FILING DATE: 2001-07-23
NUMBER OF SEQ ID NOS: 742
SOFTWARE: Patentin version 3.1
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APPLICANT: Murry, Lynn E.
TITLE OF INVENTION: HUMAN ANGIOPOIETIN
FILE REFERENCE: PC-0048 CIP
CURRENT APPLICATION NUMBER: US/09/897,306
CURRENT FILING DATE: 2001-07-02
                                                                                                                                                                                                                    LENGTH: 77
TYPE: DNA
ORGANISM:
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US20020123054A1
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Sequence 7, Applia Patent No. US2002

cation US/10072152

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GENERAL INFORMATION APPLICANT: Ander:

APPLICANT:

lein,

Lene N.

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RESULT 6
US-09-938-842A-2733
; Sequence 2733, Application
; Sequence 2733, Application
; Sequence 2733, Application
                                                                                                               APPLICANT: Zhu, Tong

APPLICANT: Zhu, Tong

TITLE OF INVENTION: STRESS-REGULATED GENES OF TITLE OF INVENTION: SAME, AND METHODS OF USE FILE REFERENCE: SCRIP1300-3

CURRENT APPLICATION NUMBER: US/09/938,842A

CURRENT FILING DATE: 2001-08-24

PRIOR APPLICATION NUMBER: US 60/227,866

PRIOR FILING DATE: 2000-08-24

PRIOR APPLICATION NUMBER: US 60/264,647

PRIOR APPLICATION NUMBER: US 60/264,647

PRIOR APPLICATION NUMBER: US 60/300,111

PRIOR FILING DATE: 2001-06-22

NUMBER OF SEQ ID NOS: 5379
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                                                       ; TYPE: DNA; ORGANISM: Arabidopsis US-09-938-842A-2733
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CURRENT FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: US/09/198,955
PRIOR FILING DATE: 1998-11-24
PRIOR APPLICATION NUMBER: 1343/97
PRIOR FILING DATE: 1997-11-24
PRIOR APPLICATION NUMBER: 1344/97
PRIOR FILING DATE: 1997-11-24
PRIOR APPLICATION NUMBER: 60/067,249
PRIOR APPLICATION NUMBER: 60/067,249
PRIOR APPLICATION NUMBER: 60/067,240
PRIOR APPLICATION NUMBER: 60/067,240
PRIOR APPLICATION NUMBER: 09/073,684
PRIOR APPLICATION NUMBER: 09/073,684
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LENGTH: 1391
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                                                                                                    LENGTH:
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APPLICATION NUMBER: 11-02
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Similarity 93.3%;
14; Conservative
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Bjornvad, Ma
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Glad, Sanne
Kauppinen, M
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Yang Tong
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RESULT 7
US-09-892-877-80/c
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US-09-948-783-79/c
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LENGTH: 1517
TYPE: DNA
ORGANISM: Homo
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OTHER INFORMATION: r
3-09-892-877-80
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Publication No.
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Publication No.
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TITLE OF INVENTION: 97 Human secreted proteins

FILE REFERENCE: PZ028P1

CURRENT APPLICATION NUMBER: US/09/892,877

CURRENT FILING DATE: 2001-06-28
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                                                                                                 APPLICATION NUMBER: FILING DATE: 1998-0: APPLICATION NUMBER:
                                                                                                                                                       APPLICATION NUMBER: 60/085,180 FILING DATE: 1998-05-12 APPLICATION NUMBER: 60/085,927 FILING DATE: 1998-05-18 APPLICATION NUMBER: 60/085,906
                                                FILING DATE: 1998-0 APPLICATION NUMBER:
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; FEATURE:
; NAME/KEY: SITE
; LOCATION: (1146)
; OTHER INFORMATION: r
US-09-892-877-115
                                                         US-09-948-783-115/c; Sequence 115, Application US/09948783; Publication No. US20030100051A1; GENERAL INFORMATION:
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SEQ ID NO 115
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                     APPLICANT: Ruben et. al.
TITLE OF INVENTION: 97 Human secreted
FILE REFERENCE: PZ028P2
CURRENT
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TITLE OF INVENTION: 97
FILE REFERENCE: PZ028P1
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R APPLICATION NUMBER: 60/08
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FILING DATE: 1998-05-18
R OF SEQ ID NOS: 465
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APPLICATION NUMBER: US/09/948,783 FILING DATE: 2001-09-10
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Pred. No. 7.9e+02;
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Pred. No. 7.9e+02;
Mismatches 1;
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US-10-025-524-33/c
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PRIOR APPLICATION NUMBER: 60/085,922
PRIOR APPLICATION NUMBER: 60/085,922
PRIOR APPLICATION NUMBER: 60/085,921
PRIOR APPLICATION NUMBER: 60/085,921
PRIOR FILING DATE: 1998-05-18
PRIOR APPLICATION NUMBER: 60/085,923
PRIOR APPLICATION NUMBER: 60/085,923
PRIOR FILING DATE: 1998-05-18
PRIOR APPLICATION NUMBER: 60/085,925
PRIOR APPLICATION NUMBER: 60/085,925
PRIOR APPLICATION NUMBER: 60/085,928
PRIOR APPLICATION NUMBER: 60/085,928
PRIOR APPLICATION NUMBER: 60/085,928
PRIOR FILING DATE: 1998-05-18
NUMBER OF SEQ ID NOS: 465
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LOCATION: (1146)
OTHER INFORMATION:
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                                                                                                                                                                                                        APPLICANT: Gallatin, W. Michael
Kilgannon, Patrick D.
TITLE OF INVENTION: ICAM-4 Materials
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
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ON NUMBER: 60/085,093
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233 South Wacker Drive,
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1999-11-10
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93.38;
 FORM: Floppy disk
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ears Tower
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US-10-025-524-33
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CURRENT APPLICATION NUMBER: US/09/887,576
CURRENT FILING DATE: 2001-06-25
PRIOR APPLICATION NUMBER: US 60/213,848
PRIOR FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: US 60/214,087
PRIOR FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: US 60/258,692
PRIOR FILING DATE: 2000-12-29
NUMBER OF SEQ ID NOS: 875
SOFTWARE: FastSEQ for Windows Version 4.0
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Best I
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                                                                                                                                                                                                                                                                                                            Patent No.
                                                                                                                                                                                                APPLICANT:
APPLICANT:
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                                                                                                                                                     APPLICANT: Cooper, Bret
TITLE OF INVENTION: Promoters
FILE REFERENCE: 1360.001US1
                                                                                                                                                                                                                                                                                              ENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 38,659
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-474-6300
TELEFAX: 312-474-0448
TELEX: 25-3856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 07/827,689
FILING DATE: 27-JAN-1992
APPLICATION NUMBER: US 07/889,724
FILING DATE: 26-MAY-1992
APPLICATION NUMBER: US 07/894,061
FILING DATE: 05-JUN-1992
APPLICATION NUMBER: US 08/009,266
FILING DATE: 22-JAN-1993
APPLICATION NUMBER: US 08/102,852
FILING DATE: 05-AUG-1993
APPLICATION NUMBER: US 08/245,295
FILING DATE: 18-MAY-1994
APPLICATION NUMBER: US 08/485,604
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
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MATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: single
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APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/10/025,524 FILING DATE: 18-Dec-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE:
                                                                                                                                                                                                                                                                             Budworth,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear ULE TYPE: DNA (g
                                                                                                                                                                                                                  Brown, D. Chang, H. Zhu, T. Han, B.
                                                                                                                                                                                                  Wang,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
PatentIn Release #1.0,
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93.3%;
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APPLICANT: Shimkets, Richard A.

APPLICANT: Jeffers, Michael

TITLE OF INVENTION: Nucleotide Sequences and Amino Acid Sequences of

TITLE OF INVENTION: Secreted Proteins Involved in Angiogenesis

FILE REFERENCE: 15966-517 CIP1

CURRENT APPLICATION NUMBER: US/10/335,819

CURRENT FILING DATE: 2003-01-02

PRIOR APPLICATION NUMBER: US/09/658,644

PRIOR APPLICATION NUMBER: U.S.S.N. 09/150,684

PRIOR APPLICATION NUMBER: U.S.S.N. 09/150,684

PRIOR FILING DATE: 1998-09-10

NUMBER OF SEQ ID NOS: 8

TORMUMBER OF SEQ ID NOS: 8
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US-09-897-306-
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US-10-335-819-1
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APPLICANT: Hillman, CAPPLICANT: Gorgone,
APPLICANT: Pattersor
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                      SOFTWARE:
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OTHER INFORMATION: Incyte ID
09-897-306-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/897,306
CURRENT FILING DATE: 2001-07-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: MUTTY, Lynn E. TITLE OF INVENTION: HUMAN ANGIOPOIETIN FILE REFERENCE: PC-0048 CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ
SOFTWARE: PERL
                                        LENGTH: 23
TYPE: DNA
ORGANISM:
FEATURE:
NAME/KEY:
LOCATION:
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14; Conservative
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CDS
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20030104573A1
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Pred. No. 8.2e+02;
0; Mismatches 1;
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PLICANY:
PLICANT: Sherw-
PLICANT: Sherw-
PLICANT: Smith, Vic-
PPLICANT: Stewart, Timoc.
PPLICANT: Tumas, Daniel
PPLICANT: Tumas, Daniel
PPLICANT: Tumas, Daniel
PPLICANT: Watanabe, Colin K
APPLICANT: Watanabe, Colin K
APPLICANT: Watanabe, Colin K
APPLICANT: Watanabe, Colin K
APPLICANT: Zhang
TITLE OF INVENTION:
FILE REFERENCE:
CURRENT APPLICATION NUMBER
CURRENT FILING DATE: 1997-C
PRIOR APPLICATION NUMBER
PRIOR FILING DATE: 1997-PRIOR FILING DATE: 1997-PRIOR FILING DATE: 1997-PRIOR FILING DATE: 1997-PRIOR APPLICATION NUMBER
PRIOR FILING DATE: 1997-PRIOR APPLICATION NUMBER
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PRIOR APPLICATION N
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Best Local S
Matches 14
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APPLICANT:
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REFERENCE:
RET APPLICATION NUMBER: US/10/028
RENT FILING DATE: 2001-12-19
R APPLICATION NUMBER: 60/049911
R FILING DATE: 1997-06-18
R APPLICATION NUMBER: 60/056974
R FILING DATE: 1997-09-26
R APPLICATION NUMBER: 60/059113
R APPLICATION NUMBER: 60/059115
R APPLICATION NUMBER: 60/059115
R APPLICATION NUMBER: 60/059117
OR APPLICATION NUMBER: 60/059117
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OR FILING DATE: 1997-09-17
OR APPLICATION NUMBER: 60/059122
OR FILING DATE: 1997-09-17
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                                                 OR FILING DATE: 1997-09-17
OR APPLICATION NUMBER: 60/059184
OR FILING DATE: 1997-09-17
OR APPLICATION NUMBER: 60/059263
OR APPLICATION NUMBER: 60/059263
OR FILING DATE: 1997-09-18
OR APPLICATION NUMBER: 60/059352
OR FILING DATE: 1997-09-19
OR APPLICATION NUMBER: 60/05988
OR FILING DATE: 1997-09-24
OR APPLICATION NUMBER: 60/062250
OR FILING DATE: 1997-10-17
OR APPLICATION NUMBER: 60/062285
OR FILING DATE: 1997-10-17
OR APPLICATION NUMBER: 60/062287
OR APPLICATION NUMBER: 60/062814
OR APPLICATION NUMBER: 60/062816
OR FILING DATE: 1997-10-24
OR APPLICATION NUMBER: 60/062816
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14; Conservative
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Filvaroff, Ellen
Gao, Wei-Qiang
Gerritsen, Mary E
Goddard, Audrey
Goddwski, Paul J.
Gurney, Austin L.
Sherwood, Steven
Smith, Victoria
Stewart, Timothy in Tumas, Daniel
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DeForge,Laura
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         NUMBER: 1997-10
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        1997-10-24
NUMBER: 60/063082
1997-10-31
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REFERENCE AUTHORS TITLE JOURNAL COMMENT RESULT 1
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DEFINITION ACCESSION VERSION KEYWORDS SOURCE FEATURE. ORGANISM source S Contact Center I National 1111 Yat Tel: 81: Fax: 81: Email: 1 mRNA sequence.
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Oryzias latipes
Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostomi;
Acanthomorpha; Acanthopterygii; Percomorpha; Atherinomorpha;
Beloniformes; Adrianichthyidae; Oryziinae; Oryzias.
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Y., Shin-i,T., Kimura,T., Narita,T.,
EST Project in Takeda's lab
Lshed (2001) : Tadasu Shin-i For Genetic Resource Information l Institute of Genetics ta, Mishima, Shizuoka 411-8540, Japan tshini@genes.nig.ac.jp. Location/Qualifiers 1. .109 /organism="Oryzias latipes" /strain="Hd-rR" Jindo, T. and Takeda,H

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Eukaryota; Metazoa; C
Mammalia; Eutheria; F
' bases 1 to 396)
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RPCI-23-269112.TV F
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: szhao@tigr.org
Clones are derived from the mouse BAC library RPCI-23. For BAC
library availability, please contact Pieter de Jong
(pieter@dejong.med.buffalo.edu). Clones may be purchased from
BACPAC Resources (http://bacpac.med.buffalo.edu/orderingframe.htm)
or from Resea ch Genetics (info@resgen.com). BAC end page:
http://www.tigr.org/tdb/bac_ends/mouse/bac_end_intro.html
Plate: 269 row: I column: 12
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AQ931789.1
GSS.
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Other_GSSs: RPCI-23-269I12.TJ
Contact: Shaying Zhao
Department of Eukaryotic Genomics
The Institute for Genomic Research
9712 Medical Center Dr., Rockville, M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ,B., Levins,M., Mcgann,S., and Fraser,C.M.
Mouse BAC End Sequences from
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/clone="MF01SSB020H21"
/clone_lib="MF01SSB cDNA"
/sex="mixture of female and male"
/tissue_type="whole embryo"
/dev_stage="segmentation stage 20
a 33 c 22 g 22 t 1
                                                                                                                                                                                                                                                                                                                                                        /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="RPCI-23-269I12"
/clone_lib="RPCI-23"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ω
                                                                        /note="Organ: Kidney/Brain; Vector: pBACe3.6; Site_1: EcoRI; Site_2: EcoRI; Female C57BL/6J mouse kidney and/or brain genomic DNA was isolated and partially digested with a combination of EcoRI and EcoRI Methylase. Size selected DNA was cloned into the pBACe3.6 vector at the EcoRI sites. The ligation products were transformed into DH10B electrocompetent cells (BRL Life Technologies). "
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                     /sex="Female"
/lab_host="DH10B"
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Rodentia;
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nn,S., Tsegaye,G., Geer,K.,
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мо. 8e+02;
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        AQ31232:
RPCI11-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Clones are derived from the human BAC library RPCI-11. For BAC library availability, please contact Pieter de Jong (pieter@dejong.med.buffalo.edu). Clones may be purchased from BACPAC Resources (http://bacpac.med.buffalo.edu/ordering_bac.htm) or from Resear h Genetics (info@resgen.com). BAC end Web Server: http://www.htsc.washington.edu Plate: 796 row: P column: 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Mahairas GG, Wallace JC, Hood L High Throughput Sequencing Center University of Washington 401 Queen Anne Avenue North, Seattle, WA Tel: (206) 616-3618 Fax: (206) 616-3887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 473)

Mahairas,G.G., Wallace,J.C., Smith,K., Swartzell,S., Holzman,T., Keller,A., Shaker,R., Furlong,J., Young,J., Zhao,S., Adams,M.D. a
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AQ66459
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Class: BAC
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Location/Qualifiers
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89 c 95 g 155 t
                                                                                                                                                                                                                                                                                                                                                                     /vrganism="Homo sapiens"
/db_xref="taxon:9606"
/clone="Plate=796 Col=5 Row=P"
/clone_lib="RPCI-11 Human Male BAC Library"
/sex="male"
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Pred. No. 1.4e+03;
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539 bp DNA
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    A linear
genomic clone
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DEFINITION

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Best Local S
Matches 15
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                                                     TITLE
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S Adams, M.D., Rounsley, S.D., Zhao, S., Bass, S., Linher, K., Golden, K., Berry, K., Granger, D., Suh, E., Wible, C., de Jong, P. and Venter, J.C. Use of human BAC End Sequences for Sequence-Ready Map Building Other_GSSs: RPCIll-112M12.TV

Contact: Shaying Zhao, William Nierman, Mark Adams Department of Eukaryotic Genomics
The Institute for Genomic Research
9712 Medical Center Dr., Rockville, MD 20850

Tel: 301 838 0200

Fax: 301 838 0208

Email: hbe@tigr.org
Clones are derived from the human BAC library RPCI-11. For BAC library availability, please contact Pieter de Jong
(pieter@dejong.med.buffalo.edu). Clones may be purchased from BACPAC Resources (http://bacpac.med.buffalo.edu/ordering) or from Research Genetics (info@resgen.com). BAC end search page: http://www.tigr.org/tdb/humgen/bac_end_search/bac_end_search.html Seq primer: SP6
Class: BAC ends.
                                                                                                                                                                                                                                                                                                                                                                                          357
                                                                                                                                                                                                                  AZ620907

1M0453M18R Mouse 10kb plasmid UI clone UUGC1M0453M18 R, DNA seque AZ620907

AZ620907.1 GI:11743097
GSS.
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AQ312321
AQ312321.1 G
GSS.
human.
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Mammalia; E
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pec, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von and Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end replasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
Similarity 100.0%;
15; Conservative
                                                                                                                              Mus musculus
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
1 (bases 1 to 609)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                       house mouse.
                                                                                                                                                                                                                                                                                                                                                                                                        AGCCATAACATGAAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /db_xref="GDB:7542923"
/db_xref="GDB:7542923"
/db_xref="taxon:9606"
/clone="RPCI-11-112M12"
/clone_lib="RPCI-11"
/sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           മ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /cell_type="Lymphocytes"
/note="Vector: pBACe3.6;
RPCIll Human Male BAC Lil
97 c 106 g 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Metazoa;
Eutheria;
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Primates;
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                                                                                                                                                                                                                                                                    sequence
                                                                                                                                                                                                                                                                                                     dq
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                                                                                                                                                Craniata; Vertebrata;
Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Library"
173 t
                                                                                                                                                                                                                                                                                   UUGC1M library Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vertebrata;
L; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ECORI;
                                                                                  Duval,B., Hamil,C.,

., Pedersen,T., Reil

., von Niederhausern,
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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                                                   reads
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; Murinae; Mus
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                                                                                                                                                                                                                                                                                                     GSS 13-DEC-2000
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RESULT 6
BH616021/c
LOCUS
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Matches
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malayi
Unpublished (2000)
Contact: Blaxter ML
Institute of Cell, Animal
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Rm. 308
84112, Univers:
Rm. 308
84112, Univers:
Rm. 308
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Class: |
High qu
                                                                                                                                                                                               Brugia
Brugia
                                                                                                                                                                                                                                BH616021 637 bp DNA
BMBAC305H05SP6_PSU Brugia malayi Genomic
malayi genomic, DNA sequence.
BH616021
BH616021.1 GI:18380709
GSS.
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Insert
                                                                                             Onchocercidae; Brugia.

1 (bases 1 to 637)

Whitton,C., Daub,J., Ware,J., Quail,M., Hall,N.,
,J., Guiliano,D., Slatko,B. and Blaxter,M.
Genome survey sequences from the human parasitic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 100.0%; 15; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Univers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Laboratory Mouse DNA Resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                 malayi.
malayi
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)1 585 7177
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sity of Utah
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Location/Qualifiers
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Length: 10000 Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0453M18"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_
                                                                                                                                                                                 Metazoa;
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                                                                                                                                                                                                                                                                                                                                                                                                                158
                            Animal
                                                                                                                                                                                Nematoda; Chromadorea; Spirurida;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15; DB 17; Pred. No. 1.5e+03; Mismatches 0;
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                             Biology
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                                                                                                Brugia
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                                                                                                                               Foster
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JOURNAL COMMENT
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AUTHORS
TITLE
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SOURCE
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ORIGIN
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Best Local
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                                                                                                                                                                                                                                       Parasitol. Today (Regul. Ed.), 16 (10), Contact: Dame JB
Dept. of Pathobiology, College of Vete University of Florida
2015 SW 23rd Avenue, Bldg 1017, Gaines Tel: 352 392 4700
Fax: 352 392 9704
Email. Approximation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AZ572103
303PvD05 Pv
AZ572103
AZ572103.1
GSS.
                                                                                                                                                                                           Email: damej@mail.vetmed.ufl.edu
Seq primer: M13(-20) forward
Class: shotgun.
                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Alveolata; Apicompi (bases 1 to 657)
Carlton, J.M.-R. and Dame, J.B.
The Plasmodium vivax and P. Pagnil Ed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          malaria parasite P. vivax.
Plasmodium vivax
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Fax: +44 131 670 5450

Email: mark.blaxter@ed.ac.uk

Sequenced from the Brugia malayi BAC library constructed b
Whitton and Dr Mike Quail. The sequence was generated by T
Pathogen Sequencing Unit, The Sanger Institute, Cambridge,
collaboration with Mark Blaxter, ICAPB, University of Edin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Seq primer: SP6
Class: BAC ends.
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/db_xref="taxon:6279"
/clone_lib="Brugia malayi Genomic Bac Library 3"
/sex="Mixed (male and female)"
/tissue_type="whole parasite"
/dev_stage="microfilaria (L1)"
/note="Vector: pBACe3.6; Site_1: BamH I; Brugia malayi genomic DNA was partially cleaved with Sau3A I and size fractionated. 7,392 clones were generated with mean insert size ~48 kbp. The library was constructed by Claire Whitton, Blaxter Nematode Genetics Lab, University of Edinburgh, UK, and Dr Mike Quail, The Pathogen Sequencing Unit, The Sanger Centre, Cambridge, UK."

109 c 120 g 212 t
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                                                                                               /organism="Plasmodium
/strain="Salvador I (0
497-598)"
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/clone_lib="Pv MBN #30"
/dev_stage="asexual blood forms"
/lab_host="Saimiri boliviensis"
/note="Vector: pBluescript SK(+)
                                                                                                                                                           1. .657
                                                                                                                                                                           Location/Qualifiers
                                                          /db_xref="taxon:5855"
/clone_lib="Pv MBN #30"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Brugia malayi"
/strain="TRS"
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                                                                                                                 um vivax"
(Collins,
                                                                                                                                                                                                                                                                                       Gainesville,
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1.5e+03;
ches 0;
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                                                                                                                                                                                                                                                                                                                          Veterinary
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409 (2000)
     vector
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ic 3', I
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  DNA,
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  phagemid
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В

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AUTHORS
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Laboratory of Functional Genetics
Bio-technology Institute of Zhejiang University
Kaixuan Road 268#, Hangzhou, Zhejiang, P.R.China
Tel: 0086-571-86892051
Fax: 0086-571-86961525
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15; Co
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Seq primer: M13 forward primer.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
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Oryza sativa
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/tissue_type="leaf"
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                                                                                                                                                                                                                                                           /organism="Oryza sativa"
/db_xref="taxon:4530"
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GSS; genome survey sequence.
Tetraodon nigroviridis.
Tetraodon nigroviridis
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Mammalia; F
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Direct Submission
Submitted (02-AUG-2001) Asao Fujiyama, The Institute of Physical and Chemical Research (RIKEN), Genomic Sciences Center (GSC); 1-7-22 Suehiro-chou, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan (E-mail:chimpbes@gsc.riken.go.jp, URL:http://hgp.gsc.riken.go.jp/, Tel:81-45-503-9111, Fax:81-45-503-9170)
Clones are derived from the chimpanzee BAC library PTB This BAC encurated during the R&D process and may have higher chance of clone tracking errors.
                                                                                                                  Tetraodon
104K08 of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BAC end sequences
Unpublished
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/clone_lib="PTB Chimpanz
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Eutheria;
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/db_xref="taxon:9598"
/clone="PTB-133B19.R"
/sex="male"
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Primates; Catarrhini;
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                Craniata;
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  Vertebrata;
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Mammalia;
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Contact
Email: (
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Charaterization and repeat analysis of the compact genome freshwater pufferfish Tetraodon nigroviridis
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Roest-Crollius, H.,
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                                     cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL, send email to:
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                                                                                                                                            : Robert Strausberg, cgapbs-r@mail.nih.gov
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Location/Qualifiers
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/db_xref="taxon:99883"
/clone="104K08"
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Eutheria; Primates;
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                                                                                                                 Ph.D.
            from Gibco
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lt,A., Quetier,F., Sauri
                                                                                                                               Christopher Moskaluk, M.D., Ph.D.,
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                                                                                                                                                             Ph.D.
                                                                                                                                                                                                                                                   Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                      o mRNA linear EST 14-JUL-2000 sapiens cDNA clone IMAGE:3133600 3',
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 17;
1.7e+03;
                                                                                                                                                                                                        Genome Anatomy Project (CGAP),
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Saurin,W., Bernot,A.
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nigroviridis
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                                                                                                                                   Michael R.
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AA953965
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                                                                                                                                                                                                           cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Se Clone distribution: NCI-CGAP clone distribution in found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 952 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham High quality sequence stop: 147.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota;
Mammalia; E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AA953965
AA953965.1
                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1997)
Contact: Robert Strausberg, |
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopl
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy
Tumor Gene Index_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human.
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/clone="IMAGE:3133600"
/clone_lib="NCI_CGAP_Kidll"
/lab_host="DH10B"
/note="Organ. "."
                                                                                                /db_xref="taxon:9606"
/clone="IMAGE:1573306"
/clone_lib="NCI_CGAP_Kid
/tissue_type="2 pooled t
/lab_host="DH10B"
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Fatima Bonaldo.
10 c 30
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                                                                                                                                                                                    organism="Homo"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Metazoa;
Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                      Christopher Moskaluk,
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Pred. No
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Mismatches 0;
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Catarrhini;
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Contact: Shoemaker
Public Sover
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Shoemaker, R., Keim, P., A., Bolla, B., Marra, M. Wylie, T., Underwood, K., Y., Person, B., Swaller, R., Ritter, E., Kohn, S., R., Waterston, R. and F. Public Soybean EST Pro-
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AW433346
AW4333346.1 GI:
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Glycine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: est@watson.wustl.edu
This clone is available through: ResGen, Invitrogen Corp.
South Memorial Parkway Huntsville, AL 35801 For further i
call: (800)-533-4363 or contact via email: ccu@resgen.com
Insert Length: 730 Std Error: 0.00
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4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108, USA
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381 bp mRNA linear EST 03-DEC-2001 sh55h12.y1 Gm-c1015 Glycine max cDNA clone GENOME SYSTEMS CLONE ID: Gm-c1015-3744 5' similar to TR:P96167 P96167 PUTATIVE ALDOLASE.;,
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Rosidae; euros
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                xhoI; This cDNA library was constructed from mRNA isolated from mature flowers of field grown plants. The cDNA library was prepared using the Stratagene pBluescript II XR cDNA library construction kit. Complementary DNA was synthesized from mRNA using a primer consisting of a poly (dT) sequence with a XhoI restriction site. EcoRI adapters were ligated to the blunt-ended cDNA fragments followed by XhoI digestion. The cDNA fragments were directionally cloned into the EcoRI-XhoI restriction site of the pBluescript vector. The ligated cDNA fragments were transformed into XL10-Gold host cells. This library was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Soybean EST Project
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and Eco RI sites of the modified pT7T3 vector. Lib went through one round of normalization. Library constructed by Bento Soares and M. Fatima Bonaldo. 60 c 72 g 91 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Glycine max"
/db_xref="taxon:3847"
/clone="GENOME SYSTEMS CLONE ID: Gm-c1015-3744"
/clone_lib="Gm-c1015"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tissue_type="Mature flowers, field grown plants"
/lab_host="XL10-Gold"
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by Dr.
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Pred. No. 4.1e+03;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            336
Randy Shoemaker and Dr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                via email: ccu@resgen.com
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Contact: Walbot V
Department of Biological Son Stanford University
855 California Ave, Palo Airel: 650 723 2227
Fax: 650 725 8221
Email: walbot@stanford.edu
Plate: 952024 row: A colu
358
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                                                                                                                                                                                                                                                                                                                                                                                                           walbot@stanford.edu
952024 row: A col
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/db_xref="taxon:4577"
/clone_lib="952 - BMS
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/dev_stage="mixed logarithmic and
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J., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn, 
Zimmerman,J. and Ecker,J.R.
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/db_xref="taxon:3702"
/clone="SALK_093049"
/clone_lib="Arabidopsis thaliana TDNA insertion lines"
/clone_lib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
92 a 102 c 84 g 125 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ecker@salk.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           stitute Genomic Analysis Laboratory (SIGnAL) k Institute for Biological Studies Torrey Pines Road, La Jolla, CA 92037, USA 453 4100 x1752 8 558 6379
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                   Leuven, Minderbroedersstraat 12
2 (bases 1 to 218)
Delabie, J. and Tierens, A.
Evidence for clonal expansion a marginal zone B cells in the ly.
Unpublished
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(G411).
X92296
X92296.1 G
germ line;
Homo sapien
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AJ227742.1
                                                                                   Direct Submission Submitted (03-MAR-1998)
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Mammalia; Eutheria;
1 (bases 1 to 218)
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Structural analysis of the VH-D-JH segments of
IgG mAb. Evidence for somatic selection
J. Immunol. 151 (7), 3604-3616 (1993)
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8376796
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Homo sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
1 (bases 1 to 209)
                                                                                                                                                      Homo sapiens.
Homo sapiens
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/db_xref="taxon:9606"
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1. .209
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Eukaryota; Me
Mammalia; Eut
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Submitted (22-JAN-1997) Pathology, University of Leuven,
Minderbroedersstraat 12, Leuven B-3000, Belgium
Location/Qualifiers
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; Eutheria; Primates;
s 1 to 219)
A. and Delabie, J.
zone lymphoma 8-VH4
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/product="IgH variable region"
/protein_id="AAB47529.1"
/protein_id="AAB47529.1"
/db_xref="GI:1841800"
/translation="SGGSISSYYWSWIRQSPGKGLEWIGYIYYSGSTNYNPSLKSRYTISVDTSKNQFSLKLNSVIAADTAVYYCAR"
54 c 56 g 56 t
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/product="immunoglobulin heavy chain variable region"
/protein_id="CAA12720.1"
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/translation="GGSVSSYYWSWIRQPPGKGLEWIGYIFYTGSTNYNPSLKSRVTI
SLDMSTDQFSLKLSSVTAADTAVYYCAR"
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<1. .>219
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/db_xref="taxon:9606"
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/db_xref="taxon:9606"
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Homo sapiens
Homo sapiens
Eukaryota; Metazoa; (
Mammalia; Eutheria; )
' 'hases 1 to 221)
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Mammalia; E
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Evidence for family genes arthritis
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VH4-GL7).
Z75359
Z75359.1
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Submitted (03-MAR-1998) Delabie
Leuven, Minderbroedersstraat 12,
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immunoglobulin; immunoglobulin heavy chain;
superfamily; immunoglobulin variable region
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<1. .>220
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/product="immunoglobulin
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Eutheria;
1 to 220)
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/db_xref="taxon:9606"
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                                                                                                                     selected humoral immune the synovial membrane of
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Catarrhini;
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Catarrhini; Hominidae;
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Baker,B.
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.W., Deane, M., Gilleece, M.H., Johnston, D., Scarffe, J.H.
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/chromosome="14"
/clone="VH4-GL7"
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/clone_lib="VH4-GL"
/germline
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67 c 56 g 54 t
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                                       Eukaryota; Metazoa; (Mammalia; Eutheria; Formula) 1 (bases 1 to 230) Voswinkel, J., Truempe
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Homo sapiens
Eukaryota; Met
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Z75348.1
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Submitted (20-AUG-2001) Braeuninger A.,
Frankfurt, Theodor-Stern-Kai 7, 60590,
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Brauninger, A., Spieker, T., Willenbrock, K., Gaulard, P., Wacker, H. Rajewsky, K., Hansmann, M.L. and Kuppers, R. Survival and clonal expansion of mutating 'forbidden' (immunoglobulin receptor-deficient) epstein-barr virus-infected cells in angioimmunoblastic t cell lymphoma

J. Exp. Med. 194 (7), 927-940 (2001)
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Homo sapiens partial IGVH4-61 gene for immunoglobulin
variable region, isolate case2-cell141.
                                                                                                                           Z75348.1 GI:1418949
immunoglobulin; immunoglobulin heavy
superfamily; immunoglobulin variable
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  Evidence for family genes
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Mammalia; Euthería; Primates;
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/db_xref="taxon:9606"
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/gene="IGVH4-61"
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H.sapiens germline immunoglobulin
VH4-GL10).
Z75347
Z75347.1 GI:1418948
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Voswinkel, J.
Direct Submission
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                                                                /organism="Homo sapiens"
/isolate="patient 8 with rheumatoid arthritis"
/db_xref="taxon:9606"
/chromosome="14"
/clone="VH4-GL10"
/cell_type="B lymphocyte"
/tissue_type="synovial membrane"
/clone_lib="VH4-GL"
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1 23,
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/isolate="patient 8 with r
/db_xref="taxon:9606"
/chromosome="14"
/clone="VH4-GL11"
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/tissue_type="synovial mem]
/clone_lib="VH4-GL"
/germline
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School, D-66421 Homburg/Saar,
Location/Qualifiers
                                                                                                                                                                                                      school,
                                                                                                                                                                                       (14-JUN-1996) Voswinkel J., chool, D-66421 Homburg/Saar, Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                 ; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 to 234)
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Pred. No. 6.1e+02;
Mismatches 0;
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DEFINITION ACCESSION VERSION
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HSVH4GL1
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AUTHORS
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HSVH4GL9
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                                    Eukaryota; Metazoa; C
Mammalia; Eutheria; P
1 (bases 1 to 235)
Voswinkel,J., Truempe
        Gause, A.
Evidence for
family genes
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H.sapiens
VH4-GL9).
Z75361
Z75361.1
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immunoglobulin; immunoglobulin heavy
superfamily; immunoglobulin variable
                                                                                                                                                                                                                                                                                                                                                                                                                                       Direct Submission
Submitted (14-JUN-1996) Voswinkel J.,
Medical School, D-66421 Homburg/Saar,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Mammalia; Eutheria; 1 (bases 1 to 234)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15;
                                                                                          HSVH4GL1 235 bp DN H.sapiens immunoglobulin heavy chain Z75346 Z75346.1 GI:1418947 immunoglobulin heavy superfamily; immunoglobulin variable Homo sapiens.
                                                                                                                                                                                                                                                  similarity 100.0%; 15; Conservative
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Voswinkel, J
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/isolate="patient 8 with r)
/db_xref="taxon:9606"
/chromosome="14"
/clone="VH4-GL9"
/cell_type="B lymphocyte"
/tissue_type="synovial memk
/clone_lib="VH4-GL"
/germline
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                                                           Chordata;
Primates;
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Primates;
synovial membrane of a patie
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                                                            Craniata; V
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; Hominidae;
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                                                                     Euteleostomi;
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VH4-GL1).
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Hum. Path
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9635678
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Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Direct Submission
Submitted (19-APR-1996) Dunn-Walters D.K., University College
Medical School, Department of Histopathology, Rockefeller Building,
University Street, London, WC1E 6JJ, UK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HSIGH2B1 236 bp DNA linear H.sapiens immunoglobulin rearranged heavy chain g 2B1). 271777 271777 271777.1 GI:1279486 diversity region; immunoglobulin; immunoglobulin joining region; variable region.
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Submitted (14-JUN-1996) Voswinkel J., Internal Medicine I,
Medical School, D-66421 Homburg/Saar, GERMANY
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
15; Conser
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AGTTACTACTGGAGC 15
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Valters,D.K., Boursier,L., Spence valters,D.K., Boursier,L., Spence valters,D.K., Boursier,L., Spence valters,D.K., Boursier,L., Spence valters,D.K., Boursier,L., Spence valter,L., Spence valter,L., Spence valter,L., Spence valter,D.K., Spence valter,
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/isolate="patient 8 with rheumatoid arthritis"
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/clone_lib="VH4-GL"
/germline
1. 235
64 c 67 g 49 t
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arity 100.0%;
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1. .236
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/rearranged
69 c 63 g
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/chromosome="14"
/clone="2B1"
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Pred. No. 6e+02;
Mismatches
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 237)
Brezinschek, H.P., Brezinschek, R.I. and Lipsky, P.E.
Analysis of the heavy chain repertoire of human periusing single-cell polymerase chain reaction
J. Immunol. 155 (1), 190-202 (1995)
95325588
7602095
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 Brauninger,A.
Rajewsky,K.,
                                                                                                                                                                                                                                                               on 100.0%;
Similarity 100.0%;
15; Conservative C
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Submitted (05-MAY-1995) Brezinschek H.P., Department of Inter Medicine, Harold C. Simmons Arthritis Research Center, Univer of Texas Southwestern Medical Center, 5323 Harry Hines Blvd., Dallas, TX 75235-8884, USA
On Oct 4, 1996 this sequence version replaced gi:1052697.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HSBDP71RB
H.sapiens
                                      Homo sapiens
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                   variable region, AJ347131 GI:1
                                                                          human.
                                                                                      AJ347131.1 GI:19773031 IGVH gene; immunoglobulin
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Homo sapiens partial IGVH4-61 g
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x87091
x87091.1 GI:159
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/db_xref="taxon:9606"
/chromosome="14"
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/note="VH4 family g
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21464858
11581315
2 (bases 1 to 238)
Braeuninger, A.
Direct Submission
Submitted (20-AUG-2001) Braeuninger A., Pathology, Frankfurt, Theodor-Stern-Kai 7, 60590, GERMANY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Survival and clonal expansion of mutating 'forbidden' (immunoglobulin receptor-deficient) epstein-barr virus-infected cells in angioimmunoblastic t cell lymphoma
J. Exp. Med. 194 (7), 927-940 (2001)
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/protein_id="CAC87531.1"
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/product="74 g 49 t
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/isolate="case2-cell150"
/db_xref="taxon:9606"
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DR WPI; 1999 myelodysplasia neuromuscular; variable heavy Agonist antibody; megakaryocyte; plabone marrow hypop 25-AUG-1997; 04-MAR-1999 21-AUG-1998; WO9910494-A2 17-JUN-1999 (GETH) GENENTECH INC. variable sapiens 1999-204 C₩, Ca ibody; thrombopoietin receptor; TPO-R; thrombopoietin; DIC; te; platelet; immunological; hematopoietic; thrombocytopenia; hypoplasia; disseminated intravascular coagulation; anemia; sia; myelotoxic chemotherapy; leukaemia; tumour; MuSK; CDR; ar; muscular dystrophy; complementarity determining region; avy chain; variable light chain; VH; VL; ss. heavy (VH) chain CDR1 encoding DNA. 666/17. rter PJ, (first entry) 97US-0918148. 98WO-US17364. Fendly BM, Gurney AL;

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Best Local
Matches 1
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17-APR-2001;
12-SEP-2001;
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immunological or hematological disorders
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2001WO-JP03288.
2001JP-0277314.
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RESULT 3
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OS Homo sapi
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PN WO2001572
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PD 09-AUG-20
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PR 26-MAY-20
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26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
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27-SEP-2000;
04-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention describes a modified antibody comprising at least 2 heavy chain variable domains and 2 or more light chain variable domains of an antibody, and exhibits thrombopoietin (TPO) agonistic effect by causing the TPO receptor to crosslink. The antibodies are useful in preventives and/or remedies for platelet reduction-associated blood diseases, thrombocytopenia following cancer chemotherapy or leukaemia. The antibody can act as a TPO signal transduction agonist by transducing a signal into cells by crosslinking a TPO receptor to exert TPO agonism. This sequence represents thrombopoietin (TPO) agonist antibody associated polynucleotide used in the creation of the modified antibody described in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200157277-A
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2000US-0632366.

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2000US-0024263.
                                                                                                                                                                   447/52.
                                                                                            derived single exon nucleic acid probes expression in human fetal liver -
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                                                                                                                                                                                                                                                           JLAR DYNAMICS INC.
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Note: The sequence data for this patent diprinted specification, but was obtained in from WIPO at ftp.wipo.int/pub/published_position.
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26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
measuring human gene expression in a sample derived from human heat present sequence is one such probe. The probes may be used for predicting, measuring and displaying gene expression in samples derived from the human heart via microarrays. By measuring gene expression, probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human heart and vascular e.g. cardiovascular disease, hypertension, cardiac arrhythmias and congenital heart disease.

Note: The sequence data for this patent did not form part of the present of the 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the
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expression (
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labeled nucleic a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        probes;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (COPD), interstitial lung di
fibrosis, neurofibromatosis,
Niemann-Pick disease, Herman
                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pulmonary dyspand hyaline mo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pulmonary alveol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            haemosiderosis
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                                                                                                                                                           01-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                  Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12011 sequences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          array; iden
algorithmi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     algorithmically predicting at least one exon from genomic sequences he eukaryote; and (b) detecting specific hybridisation of detectably led nucleic acids from eukaryote lung mRNA, to a single exon probe, ng a fragment identical to the predicted exon, the probe is included he above mentioned microarray; assigning exons to a single exon probe, rising (a) identifying exons from genomic sequence by the method early by measuring the expression of each of the exons in several tues and/or cell types using hybridisation to a single exon coarrays having a probe with the exon, where a common pattern of exons should be assigned to a single gene; a peptide comprising one ession analysis, mentioned in the specification, or encoded by the ession analysis, and for identifying exons in a gene, particularly ghuman lung derived mRNA and for the study of lung diseases as asthma, lung cancer, chronic obstructive pulmonary disease (ILD), familial idiopathic pulmonary osis, neurofibromatosis, tuberous sclerosis, Gaucher's disease, lonary alveolar proteinosis, Karagener syndrome, sarcoidosis, pulmonary nosiderosis, primary ciliary dyskinesis, pulmonary hypertension hyaline membrane disease. The present sequence is a single exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   exons
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array; identifying
                                                               sapiens
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as asthma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             open
                                                                                                                                                                                                                                                                                                       82
                                                                                                                              foetal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ф
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15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the novel set of probes .... acid expressed in the human lung; measuring the acid expressed in the human lung; measuring to contacting the derived from from human lung, comprising (a) contacting the ction of detectably labeled nucleic acids derived from ind (b) measuring the label detectably bound to each and (b) measuring the label detectably bound to each ind (b) measuring exons in a eukaryotic genome, comprising the label detectably bound to each indicate the second from genome.
                                                                                                                                                                                                                                                                                                                                     \vdash
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                                                                                              foetal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Also
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  expression in human lung samples
                                                                                                                                                                                                                                                                                                                                     ACTACTGGAGC 15
                                                                                                                                                                                                                                                                                                                                                                    onservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ading frame of the invention uence data for this patent d specification, but was olly from WIPO at
                                                                                                                            liver single
                                                                                                                                                            (first
                                                                                                                                                                                                                          dard;
                                                                                                                                                                                                                                                                                                       ACTACTGGAGC
                                                                                                                                                                                                                                                                                                                                                                                    arity
                                                                                                                                                                                                                                                                                                                                                                                                                                                               pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ncluded are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 relates to a spatially-addressable set of single exon robes for measuring gene expression in a sample derived gromprising single exon nucleic acid probes having one acid sequences mentioned in the specification, or their
                                                                                              liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                    5
                                                                                                                                                                                                                          DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   A; 47 C; 37 G; 36 T; 0 other;
                                                                                                                                                           entry)
                                                                                                                                                                                                                                                                                                                                                                                 100.0%;
                                                                                              gene expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          e a microarray comprising the novel set of probes which hybridise at high stringency to a n the human lung; measuring gene expression in a n lung, comprising (a) contacting the array with
                                                                                                                                                                                                                          281 BP.
                                                                                                                                                                                                                                                                                                       6
8
                                                                                                                             exon nucleic
                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the invention
                                                                                                                                                                                                                                                                                                                                                                                  Score 15; DB Pred. No. 81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       reading frames
                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                was obtained
                                                                                              single exon nucleic
                                                                                                                            acid probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              did not
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                                                                                                                                                                                                                                                                                                                                                                                                   24;
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                                                                                                                                                                                                                                                                                                                                                                                                  Length 171;
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                                                                                                                              #18468.
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ch probe of
                                                                                               acid probe; ss.
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ABA36928
                                                                                                                                                                                                                                  RESULT 11
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Best Local S
Matches 15
      04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
                                                                                                                                                                                                                                                                                                                                fetal liver. The present sequence is a single exon probe of the invention.

Note: The sequence data for this patent did not form printed specification, but was obtained in electronic from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                     measuring human gene expression in a sample derived from human foetal liver. The single exon nucleic acid probes may be used for predicting measuring and displaying gene expression in samples derived from human fetal liver. The present sequence is a single exon nucleic acid
                                                                                                                                                                                                                  ABA36928
                                                                                                                                                                                                                                                                                                                                                                                                                      Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                    analyzing
                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
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                                                                                                     WO200157274-A2
                                                                                                                                    Human; gene expression; h cardiovascular disease; h congenital heart disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MOLE-)
                                                                                                                                                                    Probe
                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JAN-2001;
                                                                      30-JAN-2001;
                                                                                                                                                                                    23-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                     invention relates to a single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SG,
                                                                                                                    sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001-483447
                                                                                                                                                                                                                                                                                                                                                                                                                      4;
                                                                                                                                                                    #15394
                                                                                                                                                                                                                                                                                                                                                                                                                                           genome-derived
                                                                                                                                                                                                                                                                                         . Similarity
15; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULAR DYNAMICS
                                                                                                                                                                                                                                                        AGTTACTACTGGAGC
                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 18468; 639pp +
                                                                                                                                                                                                                                                                                                                        281 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                    gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hanzel
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2000US-0207456.

2000US-0608408.

2000US-0632366.

2000US-0234687.

2000US-0236359.

2000GB-0024263.
      2000US-0180312.

2000US-0207456.

2000US-0608408.

2000US-0632366.

2000US-0234687.

2000US-0236359.

2000GB-0024263.
                                                                                                                                                                                                                                                                                         Conservative
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                                                                      2001WO-US00666
                                                                                                                                                                                                                                                                                                                                                                                                                                    expression i
                                                                                                                                                                   gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DK,
                                                                                                                                                                                                                                                                                                                        61 A; 81 C;
                                                                                                                                                                                                                  DNA; . 281 BP
                                                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                               100.0%;
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                                                                                                                                                                   expression
                                                                                                                                                    heart;
                                                                                                                                            hypertension;
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                                                                                                                                                                                                                                                                                                                        84
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                                                                                                                                            microarray; vascular
ension; cardiac arrhy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rank
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                                                                                                                                                                                                                                                                                        ed. No. 84;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleic
an fetal
                                                                                                                                                                   analysis
                                                                                                                                                                                                                                                                                                                                                                                                      exon nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                    sequence listing;
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                                                                                                                                                                                                                                                                                                       15;
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                                                                                                                                                                                                                                                                                                       DB
                                                                                                                                                                                                                                                                                                                                                                                                                                     acid probes
liver -
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                                                                                                                                                                                                                                                                                                                        other;
                                                                                                                                                                                                                                                                                                       22;
                                                                                                                                                                   human
                                                                                                                                            scular system;
arrhythmia;
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                                                                                                                                                                                                                                                                                                                                                                                                      acid
                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                   heart
                                                                                                                                                                                                                                                                                                                                                                                             probe
from |
                                                                                                                                                                                                                                                                                                                                                                                                                    English
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                                                                                                                                                                                                                                                                                                                                                                                                                                            useful
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                                                                                                                                                                                                                                                                                                        281;
                                                                                                                                                                   cell
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                                                                                                                                                                   sample
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RESULT 1
AAK18381
ID AAH
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Best Local
Matches
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                                                                                                   04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
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hearts
single
brains
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present sequence is one such probe. The probes may be used for predicting, measuring and displaying gene expression in samples derived from the human heart via microarrays. By measuring gene expression, the probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human heart and vascular systems. Congenital heart disease, hypertension, cardiac arrhythmias and congenital heart disease.
                                                                                                                                                                                                                                                                                     microarray;
epilepsy; c
                                                                                                                                                                                                                    09-AUG-2001
                                                                                                                                                                                                                                                                                                           Human;
                                                                                                                                                                                                                                                                                                                                                                              AAK18383;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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                                                                               (MOLE-)
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83
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                                                       SG,
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                                 2001-483
                                                                                                                                                                                                                                                                                                                                  brain
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-
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AGTTA
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                                                                                                                                                                                                                                                                                      cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human gene
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                                                       Hanzel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       equence data for this patent did not form part of the printed on, but was obtained in electronic format directly from WIPO int/pub/published_pct_sequences.
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                                                                                                   2000US-0180312.

2000US-0207456.

2000US-0608408.

2000US-0632366.

2000US-0234687.

2000US-0236359.

2000GB-0024263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
larity 100.0%;
Conservative 0
                                                                                                                                                                                                                                                                                                 expressed exon; gene expression analysis; probe;
lzheimer's disease; multiple sclerosis; schizophrenia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP;
                                 446/52
                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACTACTGGAGC
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            acleic
                                                                               JLAR DYNAMICS INC
                                                                                                                                                                                              2001WO-US00667.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61
                                                        DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ition relates to single exon nucleic acid probes for gene expression in a sample derived from human heart. is one such probe. The probes may be used for
                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
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            acid
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                                                        Chen
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          probes
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                                                        Rank
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          analyzing gene
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                                                                                probes
bone_ma
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microarray;
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                                                                               which are derived from genomic sequences expressed in the arrow. They can be used to measure gene expression in bone
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2000US-0207456.

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                                            on provides a mixed from genomic sequences expression be used to measure gene expression be the improved diagnosis and enable the improved diagnosis and
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21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
                                                                                            The present invention relates to human single exon nucleic acid probes (SENP). The present sequence is one such probe. The SENPs are derived from human HeLa cells. The SENPs can be used to produce a single exon microarray, which can be used for measuring human gene expression in a sample derived from human cervical epithelial cells. By measuring gene expression, the probes are therefore useful in grading and/or staging of diseases of the cervix, notably cervical cancer.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
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2000US-0608408.
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                                                                                                                                                                                                          ID No 14743; 487pp;
                                                                                                                                                                                                                             erived single exon nucleic acid probes useful for expression in human cervical epithelial cells -
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04-OCT-2000;
                                                                                                          The present invention relates to single exon nucleic acid probes (SENP). The present sequence is one such probe. The probes are useful for producing a microarray for predicting, measuring and displaying gene expression in samples derived from human placenta. The probes are useful for antenatal diagnosis of human genetic disorders.
                                                                                                                                                               Claim
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/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
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US-09-042-353-400
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Sequence 9, Appli	Sequence 9, Appli	Sequence 23, Appl	Sequence 37, Appl	Sequence 37, Appl	Sequence 37, Appl	Sequence 37, Appl		Sequence 2, Appli	Sequence 268, App	ω	Sequence 19, Appl		Sequence 7, Appli	Sequence 34, Appl	Sequence 31, Appl	Sequence 219, App	Sequence 419, App

ALIGNMENTS

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RESULT 1
US-08-918-148-42
US-08-918-148-42
Sequence 42, Application US/08918148A
Patent No. 6342220
GENERAL INFORMATION:
APPLICANT: Adams, Camellia
APPLICANT: W.
APPLICANT: Carter, Paul J.
APPLICANT: Fendly, Brian M.
APPLICANT: Gurney, Austin L.
TITLE OF INVENTION: Agonist Antibodies
FILE REFERENCE: P0979
CURRENT APPLICATION NUMBER: US/08/918,148A
CURRENT FILING DATE: 1997-08-25
NUMBER OF SEQ ID NOS: 79
SEQ ID NO 42
LENGTH: 15
TYPE: DNA
ORGANISM: artificial
FEATURE:
NAME/KEY: 12E10scfv VH CDR1
LOCATION: 1-15
OTHER INFORMATION:
US-08-918-148-42
                                                                              RESULT 2
US-08-652-816A-20
; Sequence 20, Application US/08652816A
; Patent No. 5872215
; GENERAL INFORMATION:
   APPLICANT: Osbourn, JK
; APPLICANT: McCafferty, JG
; APPLICANT: McCafferty, JG
; TITLE OF INVENTION: Specific binding members, materials and
; TITLE OF SEQUENCES: 53
; NUMBER OF SEQUENCES: 53
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CORRESPONDENCE
ADDRESSEE:
STREET: 63C
CITY: Chica
STATE: Illi
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Marshall, O'Toole, Gerstein, Murray &
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; Sequence 59, Application US/08545809A; Patent No. 6096878; GENERAL INFORMATION:
APPLICANT: Honjo, Tasuku; APPLICANT: Matsuda, Fumihiko
TITLE OF INVENTION: HUMAN IMMUNOGLO; TITLE OF: INVENTION: SEGMENTS AND DN.
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APPLICATION NUMBER: GB 9525004.9
FILING DATE: 07-DEC-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9610824.6
FILING DATE: 23-MAY-1996
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-474-6300
INFORMATION FOR SEQ ID NO: 20:
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PRIOR APPLICATION NUMBER: GB 920
FILING DATE: 23-SEP-1992
PRIOR APPLICATION
                                                       NUMBER OF SEQUENCES: 145
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson,
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: PCT/GB92/02240 FILING DATE: 02-DEC-1992 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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ATTORNEY/AGENT INFORMATION:
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FILING DATE: 02-DEC-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9
FILING DATE: 24-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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FILING DATE: 02-DEC-1991
PRIOR APPLICATION DATA:
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STRANDEDNESS: double
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CITY: 1
COUNTRY: US
ZIP: 02110-2804
                               STATE:
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REGISTRATION NUMBER: TEFERENCE/DOCKET NUMBER
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01-JUN-1994
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RESULT 4
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                                                  APPLICATION
FILING DATE:
PRIOR APPLICAT
APPLICATION
FILING DATE:
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APPLICANT: Honjo,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: line MOLECULE TYPE: GORIGINAL SOURCE:
                                                                                                                                                                ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                       ATTORNEY/AGENT
                                                                                                                  SOFTWARE: F
                                                                                                                                                                                                                                                                                APPLICANT: Matsuda, Fumihiko
TITLE OF INVENTION: HUMAN IMMUNOGLOBULIN VH GENE
TITLE OF INVENTION: SEGMENTS AND DNA FRAGMENTS CONTAINING THE SAME
NUMBER OF SEQUENCES: 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 29,066
REFERENCE/DOCKET NUMBER: 06!
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-8906
TELEFAX: 617-542-8906
                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
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NAME: Freeman, John W.
REGISTRATION NUMBER: 29,
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APPLICATION NUMBER: PCT/
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                                                                                                                                                                                                        STATE: N
COUNTRY:
                            NAME:
                                                                                                                                          COMPUTER: :
                                                                                                                                                                                                                                  CITY:
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TYPE: r
               REGISTRATION
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OPERATING
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                                                                E: 27-MAR-1996
ATION DATA:
N NUMBER: PCT/J
    OCKET
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78
                nan, John W.
N NUMBER: 2
                                                                                                                FastSEQ for Windows
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                                        INFORMATION:
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5 Franklin Street
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29,066
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RESULT 5
US-08-545-809A-61
Caruence 61, App
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Best Local Similarity
Matches 15; Conser
                                                                                              INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1212 harm
                                                                                                                                                           APPLICATION NUMBER: US/08/545,809A FILING DATE: 27-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/JP93/00603
FILING DATE: 10-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: Freeman, John W.
REGISTRATION NUMBER: 29,066
REFERENCE/DOCKET NUMBER: 06501/00400:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-8906
TELEFAX: 617-542-8906
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INFORMATION FOR SEQ
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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TELEPHONE: 617-542-5070
TELEFAX: 617-542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Matsuda, Fumihiko
TITLE OF INVENTION: HUMAN IMMUNOGLOBULIN VH GENE
TITLE OF INVENTION: SEGMENTS AND DNA FRAGMENTS CONTAINING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                        MOLECULE TYPE: (
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
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ZIP: 02110-2804
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                            ORGANISM:
                                                                                 STRANDEDNESS:
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               TYPE:
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Homo sapiens human lymphc CGM1
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la, Fumihiko
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RESULT 6
US-08-477-553A-3
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No. 5919910
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                           RESULT 7
US-08-477-553A-
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        Sequence 6, Patent No.
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                                                                                                                                          Query Match
GENERAL
                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 0073
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-6620
TELEFAX: (703) 836-2021
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                            TELEFAX: (
                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 07-JUN-19
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 23-JUNE-1
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                                                                                                                                                                  TOPOLOGY: 1
MOLECULE TYPE:
8-477-553A-3
                                                                                                                                                                                                                                                                                                             APPLICATION
FILING DATE:
ATTORNEY/AGENT
NAME: Meuth
                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: FILING DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 536 PRIOR APPLICATION DATA:
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CITY: ,
STATE:
                                                                                                                               Local
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          6, Application
5. 5919910
INFORMATION:
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                                                                                                                    93.3%; Score 14; DB 2; Similarity 100.0%; Pred. No. 17; 14; Conservative 0; Mismatches
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EDNESS: sing
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1th, Donna M.
ON NUMBER: 36
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: 23-JUNE-1992
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). Box 1404
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                    US/08477553A
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APPLICANT:

HUGHES-JONES,

Nevin C

NUMBER:

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RESULT 8
US-09-042-353-397
; Sequence 397, Application US/09042353
; Patent No. 6255458
; GENERAL INFORMATION:
; APPLICANT: Lonberg, Nils
; APPLICANT: Kav. Robert M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (703) 000 TELEFAX: (703) 000 FOR SEQ ID NO: 6 SEQUENCE CHARACTERISTICS: TENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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MOLECULE TYPE:
08-477-553A-6
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APPLICATION NUMBER: US/08,
FILING DATE: 07-JUN-1995
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
                                                                                                                                                                   APPLICANT: Lonberg, Nils APPLICANT: Kay, Robert M. TITLE OF INVENTION: Transgrittle OF INVENTION: Production NUMBER OF SEQUENCES: 421

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 23-JUNE-
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GI
FILING DATE: 13-NOV-1
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          ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE MEDIUM TYPE: FI
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 CURRENT
                                                                                                            STREET:
CITY: :
STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Meuth, Donna M.
REGISTRATION NUMBER: 36,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC
OPERATING SYSTEM:
SOFTWARE: Patent
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ZIP: 22313-1404
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1 Similarity 100.0%;
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APPLICATION DATA:
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Alexandria
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                                                                                                                California
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(703) 836-2021
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linear
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O. Box 1404
                                                                                                                             Francisco
                                                                                                                                          Townsend and Townsend
Townsend Center, F
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                                            MOLECULE 19-042-353-3
                                                                                                                                    REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
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                                                                                                            FORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER:
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FILING DATE
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NATION DATA:
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10-OCT-1996
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03-DEC-1993
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13-MAR-1998
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18-NOV-1993
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23-JUN-1992
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18-MAR-1992
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02-DEC-1996
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ER: 014643-009040US
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RESULT 9
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APPLICATION NUMBER: US
FILING DATE: 10-OCT-19
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 10-OCT-19
PRIOR APPLICATION DATA:
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APPLICATION NUMBER: US 07/810,279
FILING DATE: 17-DEC-1991
PRIOR APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and
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TITLE OF INVENTION: Transgenic
TITLE OF INVENTION: Producing I
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APPLICATION NUMBER:
FILING DATE: 03-DEC-
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APPLICATION NUMBER:
FILING DATE: 22-JUL-
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               APPLICATION NUMBER: FILING DATE: 10-OCT-PRIOR APPLICATION DATA:
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CITY: S
STATE:
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California
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Kay, Robert M.
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Floppy disk
                                                                                                                                       JMBER: US 08/544,404
10-OCT-1995
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09-MAR-1994
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Heterologous Antibodies
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US-08-758-417A-247
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Best Local
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TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 400:
SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: US/08/758,417A
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APPLICATION NUMBER: US 08/728,463
FILING DATE: 10-OCT-1996
APPLICATION NUMBER: US 08/544,404
FILING DATE: 10-OCT-1995
APPLICATION NUMBER: US 08/352,322
FILING DATE: 07-DEC-1994
APPLICATION NUMBER: US 08/209,741
FILING DATE: 09-MAR-1994
APPLICATION NUMBER: US 08/165,699
FILING DATE: 10-DEC-1993
APPLICATION NUMBER: US 08/161,739
FILING DATE: 03-DEC-1993
APPLICATION NUMBER: US 08/155,301
FILING DATE: 18-NOV-1993
APPLICATION NUMBER: US 08/096,762
FILING DATE: 22-JUL-1993
APPLICATION NUMBER: US 08/053,131
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CLASSIFICATION: <Unknown>
RAPPLICATION DATA:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CITY: San Francisco
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Kay, Robert M.

INVENTION: Transgenic No. 6300129-Human Animals for Producing Heterologous Antibodies
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NUMBER: 014643-009040US
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Pred. No.
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US-08-758-417A-250/c; Sequence 250, Appl
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Best Local S
Matches 14
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: CORRESPONDENCE ADDRE
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Kay, Robert M.
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SEQUENCE DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
MATION FOR SEQ ID NO: 247:
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APPLICATION NUMBER: US 0
FILING DATE: 16-DEC-1992
ATTORNEY/AGENT INFORMATION:
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FILING DATE: 07-DEC-1994
APPLICATION NUMBER: US 08/209,741
FILING DATE: 09-MAR-1994
APPLICATION NUMBER: US 08/165,699
FILING DATE: 10-DEC-1993
APPLICATION NUMBER: US 08/161,739
FILING DATE: 03-DEC-1993
APPLICATION NUMBER: US 08/155,301
FILING DATE: 18-NOV-1993
APPLICATION NUMBER: US 08/096,762
FILING DATE: 22-JUL-1993
APPLICATION NUMBER: US 08/053,131
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CITY: San Francisco
STATE: California
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ZIP: 94111-
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NUMBER: 014643-009030US
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Mismatches
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Heterologous Antibodie
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US-08-758-417A-250
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Patent No
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APPLICATION NUMBER: US/OF FILING DATE: 13-MAR-1998
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US OF THE PRIOR NUMBER: US O
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ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy d
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PRIOR APPLICATION
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FILING DATE:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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TITLE OF INVENTION:
TITLE OF INVENTION:
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CITY:
APPLICATION FILING DATE:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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o. 6255458
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JENCE CHARACTERISTICS:
LENGTH: 43 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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REGISTRATION NUMBER: 41,303
REFERENCE/DOCKET NUMBER: 014643-009030US
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APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
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                                                                                                                                                                                                                          MBER: US 07/904,068
23-JUN-1992
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18-MAR-1992
                                                                                                                                                        MBER: US 07/990,860
16-DEC-1992
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13-MAR-1998
   22-JUL-1993
                                                                              MBER: US 08/053,131
26-APR-1993
                                                                                                                                                                                                                                                                                                                                                                                             17-DEC-1991
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NE: (415) 576-0200
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                        us 08/096,762
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RESULT 13
US-08-758-417A-412
; Sequence 412, Application US/08758417A
; Patent No. 6300129
; GENERAL INFORMATION:
APPLICANT: Lonberg, Nils
Kay, Robert M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; MOLECULE TYPE: US-09-042-353-148
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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FILING DATE: 10-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US96/16433
FILING DATE: 10-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/758,417
FILING DATE: 02-DEC-1996
PRIOR APPLICATION NUMBER: WO PCT/US97/21803
FILING DATE: 01-DEC-1997
ATTORNEY/ACENT INCOMMETCH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 148:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/352, FILING DATE: 07-DEC-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/544, FILING DATE: 10-OCT-1995
PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
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APPLICATION NUMBER:
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APPLICATION NUMBER:
FILING DATE: 18-NOV-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
                                                                                         NUMBER OF SEQUENCES: 417
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and
STREET: Two Embarcadero
CITY: San Francisco
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
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ER: 014643-009040US
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Pred. No.
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Center, E
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                                                                                                                     l and Crew LLP
Eighth Floor
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US-09-042-353-149
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Best Local :
                                                                                                                                                                                                                                                                                                           equence 149, App
atent No. 625545
GENERAL INFORMAT
                                                                                         NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRES
ADDRESSEE: Townsen
STREET: Two Embarc
CITY: San Francisc
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                                                                                                                                                                                                               APPLICANT:
TITLE OF IN
  COMPUTER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Serafini, Andrew T.
REGISTRATION NUMBER: 41,303
REFERENCE/DOCKET NUMBER: 014643-009030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
TORMATION FOR SEQUENCE CONTACTOR ATTOMATION FOR SEQUENCE CONTACTOR A
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FILING DATE: 10-OCT-1996
APPLICATION NUMBER: US 08/544,404
FILING DATE: 10-OCT-1995
APPLICATION NUMBER: US 08/52,322
FILING DATE: 07-DEC-1994
APPLICATION NUMBER: US 08/209,741
FILING DATE: 09-MAR-1994
APPLICATION NUMBER: US 08/165,699
FILING DATE: 10-DEC-1993
APPLICATION NUMBER: US 08/161,739
FILING DATE: 03-DEC-1993
APPLICATION NUMBER: US 08/155,301
FILING DATE: 18-NOV-1993
APPLICATION NUMBER: US 08/096,762
FILING DATE: 22-JUL-1993
APPLICATION NUMBER: US 08/053,131
FILING DATE: 26-APR-1993
APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
ATTORNEY/AGENT INFORMATION:
NAME: Serafini Andrew T
                                                STATE: C
                           ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn R CURRENT APPLICATION DATA:
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14; Conserv
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                INVENTION:
 READABLE FORM:
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TOPOLOGY: linear
TULE TYPE: DNA (genomic)
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LENGTH: 243 base pairs
TYPE: nucleic acid
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                                                                                                                                                                                                                                     Kay, Robert M. ENTION: Trans
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                                                                                                                                            E ADDRESS:
Townsend a
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                                                                                                                     Townsend and Crew LLP Center, Eighth Floor
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Floppy disk

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US-09
                                                                                              TELEFAX: (415) 576-030 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: UFILING DATE: 03-DEC-1 PRIOR APPLICATION DATA: APPLICATION NUMBER: UFILING DATE: 10-DEC-1
MOLECULE TYPE: -042-353-149
                                                                                                                                                                 NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                    FILING DATE: 01-DEC-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 10-DEC-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/209,741
FILING DATE: 09-MAR-1994
PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/053,131
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APPLICATION NUMBER: US 08/728,463
FILING DATE: 10-OCT-1996
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APPLICATION NUMBER:
FILING DATE: 17-DE
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                                TOPOLOGY:
                                                  STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 62-DF
                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: WO PERFILING DATE: 10-OCT-1996
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                                                                                                                                                     TELEPHONE:
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COMPUTER: I
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                                                                                                                  : (415) 576-0200
(415) 576-0300
OR SEQ ID NO: 149
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02-DEC-1996
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10-OCT-1995
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18-NOV-1993
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17-DEC-1991
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23-JUN-1992
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18-MAR-1992
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13-MAR-1998
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ER: 014643-009040US
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Query

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RESULT 15
US-08-758-417A-413
Caruence 413, Ap
US-08-758-417A-4
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Patent No
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                                                                                                                                    INFORMATION FOR
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                   FILING DATE: 18-NOV-1993
APPLICATION NUMBER: US 08/096,762
FILING DATE: 22-JUL-1993
APPLICATION NUMBER: US 08/053,131
FILING DATE: 26-APR-1993
APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
ATTORNEY/AGENT INFORMATION:
              MOLECULE
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: PC
SOFTWARE: Patentin R
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                     SEQUENCE CI
                                                                                                                                                                                      TELECOMMUNICATION
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
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                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 08/728,463
FILING DATE: 10-OCT-1996
APPLICATION NUMBER: US 08/544,404
FILING DATE: 10-OCT-1995
APPLICATION NUMBER: US 08/352,322
FILING DATE: 07-DEC-1994
APPLICATION NUMBER: US 08/209,741
FILING DATE: 09-MAR-1994
APPLICATION NUMBER: US 08/165,699
FILING DATE: 10-DEC-1993
APPLICATION NUMBER: US 08/161,739
FILING DATE: 03-DEC-1993
APPLICATION NUMBER: US 08/155,301
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                                                                                                                              NAME: Serafini, Andrew T.
REGISTRATION NUMBER: 41,303
REFERENCE/DOCKET NUMBER: 014643-009030US
COMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
N FOR SEQ ID NO: 413:
                                                            LENGTH: 282 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/758,417A FILING DATE: 02-Dec-1996 CLASSIFICATION: <Unknown>
                                                  TOPOLO
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Kay, Robert M.
INVENTION: Transc
           DGY: linear
YPE: DNA (genomic)
ESCRIPTION: SEQ ID
                                                                                                                    HARACTERISTICS:
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California
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              NO:
              413:
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Best Local Similarity 100.0%; Pred. No. 25; Matches 14; Conservative 0; Mismatches	0;	Indels	0;	Gaps	0;
Qy 2 GTTACTACTGGAGC 15 Db 44 GTTACTACTGGAGC 57				-	
Search completed: June 26, 2003, 04:52:44 Job time : 25.2 secs					
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/cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
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/cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
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/cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US10_PUBCOMB.seq:*
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US-10-067-800-67

US-09-918-995-16699

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Sequence 22522, A
Sequence 22248, A
Sequence 28401, A
Sequence 67, Appl
Sequence 16699, A
Sequence 16650, A
Sequence 5474, Ap
Sequence 5760, Ap
Sequence 1157, Ap
Sequence 1157, Ap
Sequence 1157, Ap
Sequence 15, Appl
Sequence 15, Appl
Sequence 28159, A
Sequence 7, Appli
Sequence 3704, Ap
Sequence 3704, Ap
Sequence 31, Appli
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	Sequence 1, Appli	Sequence 9491, Ap	Sequence 9165, Ap	e 3, A	Sequence 192, App		e 9902	963,	\v	143	1.1	59, App	w	32297	e 121,	9777, P	e 74	e 19	19,	e 71,	e 81,	σ	e 1	Sequence 14715, A	ســـز	Sequence 5338, Ap

ALIGNMENTS

RESULT 1 US-09-864-761-2 ; Sequence 2252 ; Patent No. US

INFORMATION:

pplication US/09864761 0048763A1

APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REPERENCE: Acondicar X-1
CURRENT EPLING DATE: 2001-05-23
PRIOR FILING DATE: 2000-05-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-04
PRIOR FILING DATE: 2000-05-05
PRIOR FILING DATE: 2000-05-06
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-06
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR PRIOR APPLICATION NUMBER: US 00/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
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PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR APPLICATION NUMBER: PCT/US01/00662
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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RESULT 2
US-09-864-761-22248
: Sequence 22248, Application U
NO. US20020048763A1
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                                                                                                                                                                                                                                                                              APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
FITTLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE
FILE OF INVENTION: GENE EXPRESSION ANALYSIS BY
FILE REFERENCE: Aeomica-X-1
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
CURRENT FILING DATE: US 60/180,312
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Best Local
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PRIOR APPLICATION NUMBER: US 60
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 09
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: US 09
PRIOR APPLICATION NUMBER: US 09
PRIOR FILING DATE: 2001-01-29
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TYPE: DNA
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R APPLICATION NUMBER: PCT/US01/00666

R FILING DATE: 2001-01-30

R APPLICATION NUMBER: PCT/US01/00667

R FILING DATE: 2001-01-30

R APPLICATION NUMBER: PCT/US01/00664

PR FILING DATE: 2001-01-30

PR FILING DATE: 2001-01-30
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R FILING DATE: 2000-02-04
R APPLICATION NUMBER: US 60
R FILING DATE: 2000-05-26
R APPLICATION NUMBER: US 09
R FILING DATE: 2000-08-03
R APPLICATION NUMBER: GB 24
R FILING DATE: 2000-10-04
R APPLICATION NUMBER: US 60
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I: EXPRESSED IN LUNG, SIGNAL = 3.5

I: EXPRESSED IN HELA, SIGNAL = 3.1

I: EXPRESSED IN HELA, SIGNAL = 28

I: EXPRESSED IN FETAL LIVER, SIGNAL = 3

I: EXPRESSED IN ADULT LIVER, SIGNAL = 3

I: EXPRESSED IN HEART, SIGNAL = 3.1

I: EXPRESSED IN BRAIN, SIGNAL = 3.1

I: EXPRESSED IN BONE MARROW, SIGNAL = 3

I: NT HIT: 918567391, EVALUE 1.60e+00

I: EST_HUMAN HIT: BE727658.1, EVALUE 1.80e-
ER: PCT/US01/
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ER: PCT/US01/
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JUE 7.80e-01
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Cenn, Sharron G.

L'LICANT: Rank, David R.

APPLICANT: Hanzel, David K.

APPLICANT: Chen, Wensheng

TITLE OF INVENTION: GENE EXPRESSION ANAL

FILE REFERENCE: Acomica-X-1

CURRENT APPLICATION NUMBER: US/09/864,761

CURRENT FILING DATE: 2001-05-23

PRIOR APPLICATION NUMBER: US 60/180,312

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR APPLICATION NUMBER: US 09/632,366

PRIOR APPLICATION NUMBER: US 09/632,366

PRIOR APPLICATION NUMBER: GB 2'

PRIOR APPLICATION NUMBER: GB 2'

RIOR APPLICATION NUMBER: GB 2'

PRIOR FILING DATE: 2000-10
                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 3
US-09-864-761-28401
; Sequence 28401, Application US/09864761
; Sequence 28401, Application US/09864761
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Best Loo
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SEQ ID NO 22248
LENGTH: 281
TYPE: DNA
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OR APPLICATION NUMBER: US 09/608,408
OR FILING DATE: 2000-06-30
OR APPLICATION NUMBER: US 09/774,203
OR FILING DATE: 2001-01-29
OR FILING DATE: 49117
OF SEQ ID NOS: 49117
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R APPLICATION NUMBER: PCT/US01/00661

R FILING DATE: 2001-01-30

R APPLICATION NUMBER: PCT/US01/00670

R FILING DATE: 2001-01-30

R APPLICATION NUMBER: US 60/234,687

PR FILING DATE: 2000-09-21
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                 APPLICATION
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APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
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INFORMATION:
-761-22248
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INFORMATION:
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15; Conservative 0
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N: EXPRESSED IN ADULT LIVER, SIGNAL = 7.6

N: EXPRESSED IN PLACENTA, SIGNAL = 4.5

N: EXPRESSED IN LUNG, SIGNAL = 3.9

N: EXPRESSED IN HEART, SIGNAL = 5.7

N: EXPRESSED IN BRAIN, SIGNAL = 4.8

N: EXPRESSED IN BONE MARROW, SIGNAL = 6.3

N: EXPRESSED IN FETAL LIVER, SIGNAL = 6.3
                 NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence Listing Engine
                               NUMBER: PCT/US01/00666
2001-01-30
2001-01-
                                                                                                                                                                                                                                                                                                                  HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC GENE EXPRESSION ANALYSIS BY MICROARRAY
                 PCT/US01/00667
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                                               APPLICANT: Roschke, Viktor
APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven, M.
APPLICANT: Ruben, Steven, M.
TITLE OF INVENTION: Human G-protein Chemokine Refille Reference: 1488.1150001
CURRENT APPLICATION NUMBER: US/10/067,800
CURRENT FILING DATE: 2002-02-08
PRIOR APPLICATION NUMBER: PCT/US01/04153
PRIOR APPLICATION NUMBER: PCT/US01/04153
PRIOR APPLICATION NUMBER: 09/779,880
PRIOR APPLICATION NUMBER: 60/297,257
PRIOR APPLICATION NUMBER: 60/297,257
PRIOR APPLICATION NUMBER: 60/310,458
PRIOR APPLICATION NUMBER: 60/310,458
PRIOR APPLICATION NUMBER: 60/310,458
PRIOR FILING DATE: 2001-08-08
PRIOR FILING DATE: 2001-10-12
PRIOR FILING DATE: 2001-10-12
PRIOR FILING DATE: 2001-11-12
PRIOR FILING DATE: 2001-12-21
NUMBER OF SEC ID NOS: 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00662
PRIOR APPLICATION NUMBER: PCT/US01/00662
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence Listing Engine (SEQ ID NO 28401)
LENGTH: 319
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Best Loc
Matches
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                     NUMBER OF SOFTWARE:
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TYPE: DNA
ORGANISM: Homo:
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-864-761-28401
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INT HIT: U03896.1, EV
EST_HUMAN HIT: AW402
SWISSPROT HIT: P0633
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UMBER: PCT/US01/00668
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P06331, EVALUE
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No.
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ALUE 0.00e+00
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0;

Query

Score

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; NUMBER OF SEQ ID NO; SOFTWARE: FastSEQ f ; SEQ ID NO 16650; LENGTH: 438; TYPE: DNA; ORGANISM: Homo sapus-09-918-995-16650
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US-09-918-995-16699
Cacuence 16699, Ap
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; NAME/KEY: misc_formation: (1)...;
; COTHER INFORMATION:
US-09-918-995-16699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Hyseq, Inc.

TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUE
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRA
FILE REFERENCE: 20411-756

CURRENT APPLICATION NUMBER: US/09/918,995

CURRENT FILING DATE: 2001-07-30
PRIOR APPLICATION NUMBER: US/09/235,076
PRIOR FILING DATE: 1999-01-20
NUMBER OF SEQ ID NOS: 38054
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 16699
LENGTH: 413
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US-09-918-995-16650
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Best I
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Publication No.
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                                                                                                                  CURRENT APPLICATION NUMBER: US/09/918,995
CURRENT FILING DATE: 2001-07-30
PRIOR APPLICATION NUMBER: US/09/235,076
PRIOR FILING DATE: 1999-01-20
                                                                                                                                                                       APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-756
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FEATURE:
NAME/KEY:
LOCATION:
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15; Conservative
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US20030073623A1
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for Windows
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Pred. No. 44;
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RESULT 7
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DR APPLICATION NUMBER: PCT/US01/00666

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FILE REFERENCE: Aeomica-X-1
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OTHER
TYPE: DNA ORGANISM: FEATURE:
                                                                 LENGTH:
                                                                 ID NO 11831
ENGTH: 475
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OR APPLICATION NUMBER: US 09/632,366

OR FILING DATE: 2000-08-03

OR APPLICATION NUMBER: GB 24263.6

OR FILING DATE: 2000-10-04

OR APPLICATION NUMBER: US 60/236,359

OR FILING DATE: 2000-09-27

OR APPLICATION NUMBER: PCT/US01/00666

OR FILING DATE: 2001-01-30

OR APPLICATION NUMBER: PCT/US01/00667
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OR APPLICATION NUMBER: PCT/US01/00662

OR FILING DATE: 2001-01-30

OR APPLICATION NUMBER: PCT/US01/00661

OR FILING DATE: 2001-01-30

OR APPLICATION NUMBER: PCT/US01/00670

OR FILING DATE: 2001-01-30

OR FILING DATE: 2000-09-21

OR APPLICATION NUMBER: US 60/234,687

OR APPLICATION NUMBER: US 09/608,408

OR FILING DATE: 2000-06-30

OR FILING DATE: 2000-06-30
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PR APPLICATION NUMBER: PCT/US01/00664

PR FILING DATE: 2001-01-30

PR APPLICATION NUMBER: PCT/US01/00669

PR FILING DATE: 2001-01-30

PR APPLICATION NUMBER: PCT/US01/00665

PR FILING DATE: 2001-01-30

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RESULT 10
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RESULT 12
US-09-864-761-3
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; OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = US-09-864-761-11831
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US-09-764-847
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LENGTH: 21470
TYPE: DNA
ORGANISM: Homo
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GENERAL INFORMATION:
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TITLE OF IN
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC009C1
CURRENT APPLICATION NUMBER: US/10/092,154
CURRENT FILING DATE: 2002-03-07
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INVENTION: Nucleic Acids, Proteins, and Antibodies
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DATE: 2001-01-17
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FILE REFERENCE: Aeomica-A -
CURRENT APPLICATION NUMBER: US/09/864,//
CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: GB 24263.6
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OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: US 60/234,687
OR FILING DATE: 2000-09-21
OR APPLICATION NUMBER: US 09/608,408
OR FILING DATE: 2000-06-30
OR APPLICATION NUMBER: US 09/774,203
OR APPLICATION NUMBER: US 09/774,203
OR FILING DATE: 2001-01-29
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OR APPLICATION NUMBER: PCT/US01/00669

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OR FILING DATE: 2001-01-30

OR APPLICATION NUMBER: PCT/US01/00662

OR APPLICATION NUMBER: PCT/US01/00661

OR APPLICATION NUMBER: PCT/US01/00661

OR APPLICATION NUMBER: PCT/US01/00661

OR APPLICATION NUMBER: PCT/US01/00670

OR APPLICATION NUMBER: PCT/US01/00670

OR APPLICATION NUMBER: US 60/234,687

OR FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00667
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 b, Application US/09879813
US20020155453A1
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28: 49117
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EXPRESSED IN ADULT LIVER, SIGNAL = 0.44

NT HIT: M99683.1, EVALUE 0.00e+00

SWISSPROT HIT: P06331, EVALUE 1.00e-26

EST_HUMAN HIT: AW403420.1, EVALUE 0.00e+00
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CURRENT FILING DATE: 2002-11-18
PRIOR APPLICATION NUMBER: 09/828,717
PRIOR FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: 09/879,813
PRIOR FILING DATE: 2001-06-11
PRIOR APPLICATION NUMBER: PCT/GB99/03358
PRIOR FILING DATE: 1999-10-08
PRIOR APPLICATION NUMBER: GB 9822104.7
PRIOR APPLICATION NUMBER: GB 9801141.3
PRIOR APPLICATION NUMBER: GB 9901141.3
PRIOR FILING DATE: 1999-01-19
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                                                           ORGANISM: Homo sapiens FEATURE:
NAME/KEY: misc_feature LOCATION: (1)..(341)
OTHER INFORMATION:
US-10-146-505-15
    Query Match
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SEQ ID NO 1
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SEQ ID NO 15
LENGTH: 341
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Publication No.
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PRIOR FILING DATE:
NUMBER OF SEQ ID NO
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ORGANISM: Homo sapid
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)..(341)
OTHER INFORMATION:
-09-879-813-15
                                                                                                                                               LENGTH: 34
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TITLE OF INVENTION: Method of Generating Diversity
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PRIOR APPLICATION NUMBER: 09/828,717

PRIOR FILING DATE: 2001-06-04

PRIOR APPLICATION NUMBER: PCT/GB99/03358

PRIOR FILING DATE: 1999-10-08

NUMBER OF SEQ ID NOS: 87
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CURRENT APPLICATION NUMBER: US/09/879,813
CHRRENT FILING DATE: 2001-06-11
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CANT: Cumbers, Sarah J.

OF INVENTION: Method of Generating Diversity

REFERENCE: 18396/2005
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UMBER: GB 9913435.5
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OR APPLICATION NUMBER: PCT/US01/00666
OR APPLICATION NUMBER: PCT/US01/00666
OR APPLICATION NUMBER: PCT/US01/00667
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OR APPLICATION NUMBER: PCT/US01/00661
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OR APPLICATION NUMBER: US 09/608,408
OR FILING DATE: 2000-09-21
OR APPLICATION NUMBER: US 09/608,408
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: US 09/774,203
OR APPLICATION NUMBER: US 09/774,203
OR FILING DATE: 2001-01-29
BER OF SEQ ID NOS: 49117
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APPLICATION NUMBER: US 09/632,366
FILING DATE: 2000-08-03
APPLICATION NUMBER: GB 24263.6
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NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Co Unpublished (1999)

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Eco RI site shown at the beginning of the sequentissue Procurement: Louis M. Staudt, M.D., Ph.D.

CDNA Library Preparation: M.B. Soares Lab

CDNA Sequencing by: M.B. Soares Lab

Clone distribution: MGC clone distribution info found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

Incoming Sequencing MGC clone distribution info found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html
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/note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: Eco RI
Constructed from size fractionated cytoplasmic mRNA
(0.5-1.5kb). Directionally cloned. Cells provided by L
M. Staudt, Ph.D. Library preparation by Maria de Fatim
Bonaldo, Ph.D. and M. Bento Soares, Ph.D."
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/db_xref="taxon:9606"
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/tissue_type="lymph"
/cell_type="germinal center E
/cell_line="MGC85"
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Brazil
Tel: +55-11-2704922
Fax: +55-11-2707001
                                                                                                                                                                                                                      l Similar
15; Cons
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AW408669 275 bp mRNA UI-HF-BM0-ads-b-08-0-UI.rl NIH_MGC_38 Homo IMAGE:3062726 5', mRNA sequence. AW408669
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(http://
021100-2
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High qualit
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Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     asimpson@ludwig.org.br
equence was derived from the FAPESP/LICR Human Cancer Genome
t. This entry can be seen in the following URL
//www.ludwig.org.br/scripts/gethtm12.p1?t1=IL5&t2=IL5-EN0085-
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                                                                                                                                                                                                                                                                                                          /note="Organ: lung_normal; Vector: puc18; Site_1: SmaI; Site_2: SmaI; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

E 1 (bases 1 to 299)

E Adams, M.D., Kerlavage, A.R., Fleischmann, R.D., Fuldner, R.A., Bult, C.J., Lee, N., Kirkness, E.F., Weinstock, K.G., Gocayne, J.D., White, O., Sutton, G., Blake, J.A., Brandon, R.C., Chiu, M.-W., Clayton, R.A., Cline, R.T., Cotton, M.D., Earle-Hughes, J., Fine, L.D., FitzGerald, L.M., FitzHugh, W.M., Fritchman, J.L., Geoghagen, N.S.M., Glodek, A., Gnehm, C.L., Hanna, M.C., Hedblom, E., HinkleJr, P.S., Kelley, J.M., Klimek, K.M., Kelley, J.C., Liu, L.-I., Marmaros, S.M., Merrick, J.M., Phillips, C.A., Ryder, S.E., Scott, J.L., Saudek, D.M., Shirley, R., Small, K.V., Spriggs, T.A., Utterback, T.R., Weidman, J.F., Li, Y., Bednarik, D.P., Cao, L., Cepeda, M.A., Coleman, T.A., Collins, E.-J.,
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Seq primer: M13 Forward.
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T27727
T27727.1 GI:60
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Contact: Robert Strausberg, Ph.D.
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Email: cgapbs-r@mail.nih.gov
Eco RI site shown at the beginning of the so
Eco RI site shown at the beginning of the so
The procurement: Louis M. Staudt, M.D.,
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National Institutes of Health, Mammalian
Unpublished (1999)
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Eutheria;
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Contact: Venter,
The Institute for
932 Clopper Rd, (932 Clopper Rd)
Tel: 3018699056
Fax: 3018699423
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,M.R., Ro
Venter,J.
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Mammalia; E
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AW405472

AW405472.1 GI:6924529
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For clone availability, additional sequence and expression
information related to this EST, please contact the TIGR Database
(tdbinfo@tdb.tigr.org)
Seq primer: M13 Reverse.
Location/Qualifiers
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Homo sapiens
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site shown at the beginning of the sequence
e Procurement: Louis M. Staudt, M.D., Ph.D.
Library Preparation: M.B. Soares Lab
Library Arrayed by: M.B. Soares Lab
equencing by: M.B. Soares Lab
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pon 83 Million Basepairs of cDNA Sequence
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for Genomic Research
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100.0%; Pred. No. 1e+03;
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Mammalia; Eutheria; Primates; Catarrhini;
1 (bases 1 to 367)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
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Email: cgapbs-r@mail.nih.gov
Eco RI site shown at the beginning of the s
Tissue Procurement: Louis M. Staudt, M.D.,
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Constructed from size fractionated cytoplasmic mRNA
(0.5-1.5kb). Directionally cloned. Cells provided by Louis
M. Staudt, Ph.D. Library preparation by Maria de Fatima
Bonaldo, Ph.D. and M. Bento Soares, Ph.D."
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Constructed from size fractionated cytoplasmic mRNA
(1.5-2.5kb). Directionally cloned. Cells provided by
M. Staudt, Ph.D. Library preparation by Maria de Fal
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      Contact: Email: c
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Eukaryota; Metazoa; Chordata; Craniata; Ve
Mammalia; Eutheria; Primates; Catarrhini;
1 (bases 1 to 459)
                                                      National Institute Unpublished (1999)
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OUI-HF-BK0-aas-f-08-0-UI.rl NIH_MGC_36 Homo sapiens cDNA clone IMAGE:3054878 5', mRNA sequence.

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IMAGE:3062322 5', mRNA sequence.
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Location/Qualifiers
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Imail: cgapbs-r@mail.nih.gov
Ico RI site shown at the beginning of the sequence
Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
cDNA Library Preparation: M.B. Soares Lab
cDNA Library Arrayed by: M.B. Soares Lab
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Clone distribution: MGC clone distribution
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Eco RI site shown at the beginning of the sequence.

Tissue Procurement: Louis M. Staudt, M.D., Ph.D.

cDNA Library Preparation: M.B. Soares Lab

cDNA Sequencing by: M.B. Soares Lab

Clone distribution: MGC clone distribution informati

found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

Seq primer: M13 Forward.
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Clone distribution: MGC clone distribution
found through the I.M.A.G.E. Consortium/LLNL
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UI-HF-BK0-aal-b-06-0-UI.rl NIH_MGC_
IMAGE:3053915 5', mRNA sequence.
AW402332
AW402332.1 GI:6921018
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National Institutes of Health, Mammalian
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Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1999)
Contact: Robert Strausberg,
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/note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: ____
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/db_xref="taxon:9606"
/clone="IMAGE:3053915"
/cell_type="germinal center
/cell_line="MGC85"
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Pred. No. 1.2e+03;
Mismatches 0;
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Contact: Rob
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AW402282
AW402282.l GI:6920968
EST.
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:3054035"
/clone_lib="NIH_MGC_36"
/tissue_type="lymph"
/cell_type="germinal center B cells"
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Institutes of Health, Mammalian Gene Collection (MGC)
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Eutheria; Primates;
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammaliar Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Eco RI site shown at the beginning of the Tissue Procurement: Louis M. Staudt, M. cDNA Library Preparation: M.B. Soares I
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Fax: 402 762 4390
Email: smith@email.marc.usda.gov
Single pass sequencing. Bases called and al: v0.980904.e. Vector identified by cross_mate and -minmatch 12 options.
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119835 MARC
AW786572
AW786572.1
EST.
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AW402602

AW402602.1 GI:6921304
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15; Conserv
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Fahrenkrug, S.C., Freking, B.A., Rohrer, G.A., Smith, T. Stone, R.T., Heaton, M.P., Grosse, W.M., Bennett, G.A., and Keele, J.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       \vdash
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Plate: 42 row: D column: 18
Seq primer: ATTTAGGTGACACTATAG.
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                                                                                                                                                                                                                 Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 548)
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                                                                                                                                                                                                                                                                                                                                                       human
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/db_xref="taxon:9823"
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/tissue_type="pooled"
/lab_host="DH10B"
/note="Vector: pCMV SPORT6; Site_1: NotI; Site_note="vector: pCMV sport6; Site_1: NotI; Site_note=
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Pred. No. 1.3e+03;
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                                  the sequence M.D., Ph.D.
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Laegreid,W.W.
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Sakyo-ku
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Phlebobranchia; Cionidae;
'hases 1 to 590)
'Kohar
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Ciona
Ciona
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Satoh, N., Satou, Y., Kohara, Y. and Shin-i,
Expressed genes in Ciona intestinalis
Unpublished (2000)
Contact: Nori Satoh
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Seq primer: M13 Forward.
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AV983832
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Sequencing by: M.B. Soares Lab
he distribution: MGC clone distribution information can
through the I.M.A.G.E. Consortium/LLNL at:
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                                                                                                                                                                                                                                                                                                                                                                            ku, Kyoto, Kyoto 606-8502, Japan
1-75-753-4081
1-75-705-1113
satoh@ascidian.zool.kyoto-u.ac.jp.
Location/Qualifiers
 100.0%;
rity 100.0%;
nservative 0;
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/db_xref="taxon:9606"
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/clone_lib="NIH_MGC_36"
/tissue_type="lymph"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .a; Metazoa; Chordata; Urochordata; Ascidiacea; anchia; Cionidae; Ciona.
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/note="Vector: pT7T3-Pac; Site_1: NotI;                                                                                                                                  /organism="Ciona intestinalis"
/db_xref="taxon:7719"
/clone="cieg28e23"
/clone_lib="Nori Satoh unpublished cDNA library,
/tissue_type="whole animal"
/dev_stage="egg"
/note="Vector: pBluescript SK"
152 c 101 g 137 t
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/cell_line="MGC85"
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Staudt, F...
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Search completed: June Job time : 753.2 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

E 1 (bases 1 to 607)

S Kim,N.S.; Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R., Ch,Kim,Y.S.

21C Frontier Korean EST Project 2001

Unpublished (2002)

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BM783015
BM783015.1 GI.1.
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K-EST0060735 S18N669761 Homo
5', mRNA sequence
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                                                                                                                                                       100.0%; Similarity 100.0%; 15; Conservative C
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Location/Qualifiers
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Plate: 2 row: E column: 06
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                                                                                                   AGTTACTACTGGAGC
                                                                                                                                                                                                                                                    /note="Organ: Stomach; Vector: pCNS; Site_1: EcoRI; Site_2: NotI; The poly (A)+ RNA was dephosphorylated with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was circularized with E. coli DNA ligase after digestion of EcoRI which site is also included in vector was converted to a DNA strand by Okayama-Berg method. The obtained cDNA vectors were used for transformation of competent cells E. coli Top10F' by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
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/clone="S18N669761-2-E06"
/clone_lib="S18N669761"
/sex="F"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     'lab_host="Top10F'"
                  26,
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3., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and
                    04:50:22
                                                                                                                                                           0;
                                                                                                                                                           Score 15; DB 14;
Pred. No. 1.3e+03;
Mismatches 0;
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